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Database
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Sequence 42, Appl	Sequence 45, Appl	Sequence 2, Appli	Sequence 25, Appl	ς	Sequence 2, Appli	•	Sequence 3, Appli		Sequence 1, Appli	Sequence 60, Appl		Sequence 18, Appl	Sequence 1, Appli	Sequence 1, Appli	Sequence 17, Appl	Sequence 17, Appl	Sequence 36, Appl

ALIGNMENTS

```
APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Sleeman, Matthew
APPLICANT: Ourist, Rene
APPLICANT: Within Strack From Skin Cells
TITLE OF INVENTION: James Greg
FILE REFERENCE: 11000.1011cl
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEQ ID NOS: 348
SOOTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 331
LENGTH: 299
TYPE: PRT
ORGANISM: Human
US-09-188-930-331
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Matches 100
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                                  AVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTS--EEGDFRHKSSFVI 310
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Sequence 2, Application US/09462270

Patent No. 6358707

GENERAL INFORMATION:

APPLICANT: SmithKline Beecham Corporation

ITLE OF INVENTION: Human F11 Antigen: A Cell Surface

TITLE OF INVENTION: Receptor Involved in Platelet Aggre

FILE REFERENCE: GH-70150US

CURRENT APPLICATION NUMBER: US/09/462,270

CURRENT FILING DATE: 2000-01-05

PRIOR APPLICATION NUMBER: 60/052,186

PRIOR APPLICATION NUMBER: 60/052,186

PRIOR FILING DATE: 1997-07-10

NUMBER: OF SEQ ID NOS: 4

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 2

LENGTH: 299
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 189
LENGTH: 299
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TYPE: PRT
ORGANISM: HOMO SAPIENS
-09-462-270-2
-09-188-930-189
                                                                                                                                                                                                                                                                                                                APPLICANT: Watson, James D.
APPLICANT: Strachan, Lorna
APPLICANT: Strachan, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Onrust, Rene
TITLE OF INVENTION: Compositions Isolated From Skin
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 1100,1011c1
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                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/188,930A CURRENT FILING DATE: 1998-11-09
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LOCATION: (247)...(247)
NAME/KEY: UNSURE
LOCATION: (289)...(289)
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                                                                                                                                  ORGANISM: Human
                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             quence 189, App
tent No. 615050
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/09188930A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25.4%; Score 415; DB 4; Length 299; 33.8%; Pred. No. 8.8e-34;
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FILE REFERENCE: GH-70150US
CURRENT APPLICATION NUMBER: US/09/462,270
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 60/052,186
PRIOR FILING DATE: 1997-07-10
NUMBER OF SEQ ID NOS: 4
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
LENGTH: 205
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                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 64; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION: APPLICANT: SmithKl
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Best Local Similarity
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TITLE OF INVENTION: Receptor Involved in Platelet Aggregation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
                            198 NSETGTLVFNA 208
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                                                                                                                                                                                                                                78 VYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVALNDRKEVDEITIELIVQV 137
                                                                                                                                                                                                                                                                                15 FILAILLCSLALGSVTVHSSEPEVRIPENNPVKLSC--AYSGFSSPRVEWKFDQGDTTRL 72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQ-DMEVYDLNIAGIIGGVLVVLIVL
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NPTTGELVEES
                                                                                          PPSKPTVNIPSSATIGNRAVLTCSEQDGSPPSEYTWFKDGIVMPTNPKSTRAFSNSSYVL
                                                                                                                                     KPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPLPTDSRANPREQNSSFHV 197
                                                                                                                                                                                       VCYNNKITASYEDRV-TFLPTGITFKSVTREDTGTYTC-MVSEEGGNSYGEVKVKLIVLV
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                                                                                                                                                                                                                                                                                                                                                                      17.3%; Score 283; DB 4; Length 205, 33.5%; Pred. No. 9.8e-21; ative 34; Mismatches 89; Indels
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33.1%; Pred. No. 1.4e-32;
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RESULT 5 US-08-597-495B-22 · Sequence 22, Application US/08597495B

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
NAME: HANSON, NO. 5712369man D.
REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: LUD 5316
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Simpson, Richard J.; Nice, Edouard; Moritz, R. L.;
APPLICANT: Catimel, B. Ji, Hong: Burgess, Anthony W.;
APPLICANT: Heath, Joan K.; White, Sara J.; Johnstone, Cameron
TITLE OF INVENTION: Colon Cell And Colon Cancer Cell
TITLE OF INVENTION: Associated Nucleic Acid Molecules, Protein And Peptides
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 319 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DC
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                    187
                                                                                                                                                                                                                                                                                                    134
                                                                                                                                                                                                                                                                                                                         132 ELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDV-----PLPTDSRA 186
                       294 EREEEDDYRQE 304
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STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                   84 --IQGDL-----AGRTDVFGKTSLRIWNVTRSDSAIYRCEVVALNDRKEVDEITI 131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 08/511,876
                                                                   RTSEEGDFRHK 305
                                                                                                                ----VGIAVGVVAALIIIGII-IYCCCCRGKDDNTEDKEDARPNREAYEEPPEQLRELSR 293
                                                                                                                                                         DLNIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDG----ESYKSPGKHDGVNYI 294
                                                                                                                                                                                                                                                 NPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAG-----AARCEGQDMEVY 238
                                                                                                                                                                                                                                                                                               RLLVLVPPSKPECGIEGETIIGNNIQLTCQSKEGSPTPQYSWKRYNILNQEQPLAQPASG 193
                                                                                                                                                                                                                                                                                                                                                                                        NYIHGELYKNRVSISNNAEQSD----ASITIDQLTMADNGTYECSVSLMSDLEGNTKSRV 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VDAISVETPQDVLRASQGKSVTLPCTY-HTSTSSREGLIQWDKLLLTHTERVVIWPFSNK 77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              49; Mismatches 123;
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Nice, Edouard; Moritz, R. L.;
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Patent No. 6391235

GENERAL INFORMATION:
GENERAL INFORMATION:
Simpson, Richard J.; Welt, Sydney; Ritter, Gerd;
Simpson, Richard J.; Nice, Edouard; Moritz, R. L.;
Catimel, B.; Ji, Hong; Burgess, Anthony W.;
Heath, Joan K.; White, Sara J.; Johnstone, Cameron
TITLE OF INVENTION: Colon Cell And Colon Cancer Cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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INFORMATION FOR SEQ ID NO:
                                             239
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                                                                                                                                                                              134 RLLVLVPPSKPECGIEGETIIGNNIQLTCQSKEGSPTPQYSWKRYNILNQEQPLAQPASG
                                                                                                                                                                                                                          132 ELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDV-----PLPTDSRA 186
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                                                                                                                                                                                                                                                                         78 NYTHGELYKNRYSISNNAEQSD----ASITIDQLTMADNGTYECSVSLMSDLEGNTKSRV
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                                                                                                                                                                                                                                                                                                                                                                                                             29 IEAVNLKSSNRNPVVHEFESVELSCIITHSQTS--DPRIEWKKIQDGQTTYVY---FDNK 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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APPLICATION NUMBER: 08/597,495
FILING DATE: 02-Feb-1996
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOTWARE: Wordperfect
CURRENT APPLICATION DATA:
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ADDRESSEE: Fulbright & Jaworski LLP
STREET: 666 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 318-3168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
----VGIAVGVVAALIIIGII-IYCCCCRGKDDNTEDKEDARPNREAYEEPPEQLRELSR 293
                                           DLNIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDG----ESYKSPGKHDGVNYI
                                                                                        QP-----VSLKNISTDTSGYYICTSSNEEGTQFCNITVAVRSPSMNVALY
                                                                                                                                   NPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAG------AARCEGQDMEVY 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Hanson, NO. 6291235man D. REGISTRATION NUMBER: 30,946 REFERENCE/DOCKET NUMBER: LUD 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/068,051A FILING DATE: 10-Dec-1998 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 319 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: New York City
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette, 3.5 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14.8%; Score 242.5; DB 4; 25.7%; Pred. No. 2.2e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Associated Nucleic Acid Molecules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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Length 319; Indels

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Gaps

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RESULT

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US-09-068-051A-32
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Patent No. 6291235
GENERAL INFORMATION:
APPLICANT: Old, Lloyd J.; Welt, Sydney; Ritter, Gerd;
Simpson, Richard J.; Nice, Edouard; Moritz, R. L.;
Catimel, B.; Ji, Hong; Burgess, Anthony W.;
Heath, Joan K.; White, Sara J.; Johnstone, Cameron
TITLE OF INVENTION: Colon Cell And Colon Cancer Cell
                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID -09-068-051A-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 32
 199
                                                                           154
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                                                                                                                                                                                                                                                                 48 SVELSCII-THSQTSDPRIEWKKIQDGQT-----TYVY---FDNKIQGDLAGRT 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/068,051A
FILING DATE: 10-Dec-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/597, 495
FILING DATE: 02-Feb-1996
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (212) 318-3168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 360 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSE: Fulbright & Jaworski LLP
STREET: 666 Fifth Avenue
STREET: New York City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                 GKTATLQCQESEGYPRPHYSW-----YRNDVPLPTDSRANPRFQNSSFHVNSETGTLVFN 207
NISTETAGYYICTSSNDVGIESCNITVAPRPPSMNIALY----AGIAGSVFVALIIIGVI 254
                                                                                                                                                                                        DVFGKTSLRIWNVTRSDSAIYRCEVVALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPV 152
                                     AVHKDDSGQYYCIASNDAGAARCE------GQDMEVYDLNIAGIIGGVLVVLIVLAVI 259
                                                                       GNNIQLTCHSAEGSPSPQYSWKSYNAQNQQRPLTQPVSGEP
                                                                                                                                                    AELSNASITIDQLTMDDNGTYECSVSLMSDQDVNAKSRVRLLVLVPPSKPDCSIQGEMVI 153
                                                                                                                                                                                                                              SVTLPCTYNTYVSDREGFIQWDKLLRSQTERVVTWNFVTKKYIYGNRYENRVR----VSND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Hanson, No. 6291235man D.
REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: LUD 5316.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (212) 752-5958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Wordperfect
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 318 amino acids
                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                           14.7%; Score 240.5; DB 4; 24.9%; Pred. No. 3.6e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Associated Nucleic Acid Molecules, Protein And Peptides
                                                                                                                                                                                                                                                                                                         37; Mismatches 122;
                                                                                                                                                                                                                                                                                                                                                                                                                          NO:
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                                                                                                                                                                                                                                                                                                                                           Length 318;
                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                         61;
                                                                         -LLLK 198
                                                                                                                                                                                                                                                                                                       Gaps
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Best Local Similarity
Thes 76; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens US-09-175-928-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-175-928-2
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APPLICANT: McCoy, John M.
APPLICANT: LaVallie, Edward R.
APPLICANT: Collins-Racie, Lisa A
APPLICANT: Evans, Cheryl
APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
                                                                                                                                                                                  Sequence 2, Application US/08928383B Patent No. 6210921 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver.
SEQ ID NO 2
LENGTH: 387
TYPE: PRT
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Patent No. 631292
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                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD,
                                                                                                                                                                                                                                                                                                                                                                                                      178 EGRDI-VPVKENFNP---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           63
             CITY: Boston
                                 STREET:
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/09175928A
                                 28 State Street
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CURRENT APPLICATION NUMBER: US/09/175,928A CURRENT FILING DATE: 1998-10-20 NUMBER OF SEQ ID NOS: 62
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APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS
FILE REFERENCE: 6006B.AJ172A
                                                                                                                                                                           APPLICANT: Robert W. Finberg, Jeffrey M. APPLICANT: and Marshall S. Horwitz TITLE OF INVENTION: CAR, A No. 6210921el TITLE OF INVENTION: Receptor NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       234 DMEVYDLNIA----GIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQ 277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        117 VVALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYR- 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68 -KK------IQDGQTTYV-YFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCE 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 FFLLLLFRGCM-----IEAVNLKSSNRNPVVHEFESVELSCIITHSQTSDPR--IEW--- 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 FWKVFLILSCLAGQVSVVQVTIPDGFVNVTVG--SNVTLICIYTTTVASREQLSIQWSFF 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VYCCCCREKD----DKDQDREDARPNRAAYQVPKKEQKEISRGREDEDDHRHE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ----IDLTSSHPEVGIIVGALIGSLVGAAIIISVVCFARNKAKAKAKE 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ---NDVPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQ 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VNNPPDFLGQNQGILNVSVLVKPSKPLCSVQGRPETGHTISLSCLSALGTPSPVYYWHKL 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HKKEMEPISIYFSQGGQAVAIGQFKDRITG-----SNDPGNASITISHMQPADSGIYICD 117
Massachusetts
: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ----TTGILVIGNLTNFEQGYYQCTAINRLGNSSCE-- 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AND POLYNUCLEOTIDES ENCODING THEM
                                                                                                                                                                                                                                             Coxsackievirus
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US-08-979-424-3
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                                                                                                                                                                                                                                                                                Sequence 3, Application US/08979424 Patent No. 5942606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 10.6
Best Local Similarity 26.4
Matches 70; Conservative
                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                          APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: VIRAL RECEPTOR PROTEIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (617)742-4214
NFORMATION FOR SEQ ID NO: 2:
                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 365 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REBERENCE/DOCKET NUMBER: DFI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/026,100
FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/928,383B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       206 FNAVHKDDSGQYYCIASNDAGAARCEGQDMEVYDLNIAGIIGGVLV-VLIVLAVITMGIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  148
                                                                                                    STREET: 31/4 FOR CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                            259 CCRK-----KRREEKYEKEVHHD 276
                                                                                                                                                                                                                                                                                                                                                                                                                             265 CAYRRGCFISSKQDGESYKSPGKHD 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  204 ----SEYSGTYSCTVRNRVGSDQCLLRLNVVPPSNKAGLIAGAIIGTLLALALIGLIIF 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           150 GSEEIGSDFKIKCEPKEGSLPLQYEWQKLSDSQKMPTSSLA--EMTSSVISVKNAS----
                                                        COUNTRY: USA
ZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93 FTSNDLKSGDASINVTNLQLSDIGTYQCKV---KKAPGVANKKIHLVVLVKPSGARCYVD 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94 V-----FGKTSLRIWNVTRSDSAIYRCEVVALNDRKEVDEITIELIVQVKPVTPVCRIP 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            35 ETAYLPCKFTLSPEDQGPLDIEWLISPADNQKVD--QVIILYSGDKIYDDYYPDLKGRVH 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47 ESVELSCIITHS-QTSDP-RIEW-----KKIQDGQTTYVYFDNKIQG----DLAGRTD 93
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CLASSIFICATION:
                                                                                                                                                  ADDRESSEE:
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                                                                                                                             E: Incyte Pharmaceuticals, 3174 Porter Dr.
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IBM Compatible
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APPLICANT: DeGregori, James
TITLE OF INVENTION: Adenoviral mediated gene tra
FILE REFERENCE: 90-98
CURRENT APPLICATION NUMBER: US/09/272,496
CURRENT FILING DATE: 1999-03-19
EARLIER APPLICATION NUMBER: US 60/092782
EARLIER FILING DATE: 1998-07-14
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                  US-09-272-496-2
; Sequence 2, Application US/09272496
; Patent No. 6245966
; ORGANISM: Homo US-09-272-496-2
                                                                                                                                                                                                                                                                                                                                                                            RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; CLONE: US-08-979-424-3
                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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Best Local Similarity 25.98
Matches 68; Conservative
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                                         TYPE: PRT
                                                          LENGTH: 365
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TELEPHONE: 650-855-0555
TELEFAX: 650-845-4166
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LENGTH: 365 amino acids
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NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                         261 RK-----KRREEKYEKEVHHD 276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          35 ETAYLPCKFTLSPEDQGPLDIEWLISPADNQKVD--QVIILYSGDKIYDDYYPDLKGRVH 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
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25.9%;
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RESULT 12
US-08-799-424-1
; Sequence 1, Application US/08979424
; Patent No. 5942606
; GENERAL INFORMATION:
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US-08-979-424-1
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Best Local Similarity
                                                                                                                                                    INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                     IMMEDIATE SOURCE:
LIBRARY: LUNGF
CLONE: 1232054
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
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APPLICANT: Corley, Neil C.
TITLE OF INVENTION: VIRAL RECEPTOR PROTEIN
                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA: APPLICATION NUMBER:
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                                                                                          TYPE: amino acid
STRANDEDNESS: single
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
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CITY: Palo Alto
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                                                                         TOPOLOGY:
                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                                                                           NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
                                                                                                                                 LENGTH:
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                                                                                                                                                                                                                                                                  PF-0405 US
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Best Local Similarity
Matches 80; Conserv
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                                                                      TELEFAX: 610-407-0701
TELEX: 846169
INFORMATION FOR SEQ ID NO:
                                                                                                                            REFERENCE/DOCKET NUMBER: GH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-407-0700
                                                                                                                                                                             APPLICATION NUMBER: 60/059,4
FILING DATE: 22-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: PRESTIA, PAUL F
REGISTRATION NUMBER: 23,031
                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: WU, SHUJIAN
APPLICANT: SWEET, RAYMOND
APPLICANT: TRUNEH, ALEMSEGED
TITLE OF INVENTION: A HUMAN LIG-1 HOMOLOG (HLIG-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                    SEQUENCE CHARACTERISTICS:
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                                                                                         TELEPHONE: 610-407-0701
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CITY: VALLEY FORGE
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                                                                                                                                                                                                                                                                                                                     FILING DATE: 08-DEC-1997
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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                                 LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSQPWEVPFVMWFFKQKEKEDQVLSYI----NGVTTSKPGVSLVYSMPSRNLSLRLEGLQE
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             amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
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O. BOX 980
                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette
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                                                                                                                                                                                                                                                              60/059,448
                                                                                                                                                                                                                                                                                                                                        US/08/986,485
                                                                       2:
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Pred. No. 4.3e-09;
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; MOLECULE TYPE: protein US-08-374-834-16
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                            TELEFAX: (914) 345-77
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Valenzuela, et al.
TITLE OF INVENTION: NOVEL TYROSINE KINASE RECEPTOR
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/095,658 FILING DATE: 21-JUL-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: UFILING DATE: 19-JAN-1 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             102 IWNVTRSDSAIYRCEVVALNDRKEVDEITIELIVQVKPVTPVCRIP---AAVPVGKTATL 158
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
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                                                       STRANDEDNESS
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                                                                                                                                                                                                                                     NAME: Cobert, Robert J.
REGISTRATION NUMBER: 36,108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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                                       TOPOLOGY:
                                                                                                     LENGTH:
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                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QCQESEGYPRPHYSWYRNDVPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYY 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        I-----DDAGVYSC--TAQNSAGSISANATLTVLE----TPSLVVPLEDRVVSVGETVAL 720
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                                                                              amino acid
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                                                                                                     869 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                             (914) 345-7721
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22.8%; Pred.
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Pred. No. 7e-08;
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US-08-644-271-29
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; MOLECULE TYPE: US-08-644-271-29
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Best Local Similarity 28.8%;
Matches 70; Conservative 2
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CLASSIFICATION: 435

PRIOR APPLICATION NUMBER: USSN 6(
APPLICATION NUMBER: USSN 6(
TOTAL DATE: 15-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                      INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS LENGTH: 869 amino acid
                                                                                                                                                                            REGISTRATION NUMBER: 36,108
REFERENCE/POCKET NUMBER: TELECOMMUNICATION INFORMATION:
TELEPHONE: 914-345-7400
                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Cobert Robert T
                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                 STRANDEDNESS
                                                                                                                                             TELEPHONE: 914 C. TELEPHONE: 914-345-7721
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                       TYPE:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 10-MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  63 -- PRIEWKK----IQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCE 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28 MIEAVNLKSSNRNPVVHEFESVELS------CIITHSQTSD------
                                  TOPOLOGY:
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                                                                       amino acid
                                                                                                                                                                                                                                               Cobert, Robert J
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'ENTION: NOVEL TYROSINE KINASE RECEPTORS
                                    unknown
                                                                                                                                                                                                                                                                                                                                                  UMBER: US/08/644,271
10-MAY-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23; Mismatches
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Search completed: September 19, 2002, 17:33:15 Job time: 233 sec
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                                                                                                        175 RNDVPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQD 234
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                                                                                            156 KGDSPLRENSRI------AVLESGSLRIHNVQKEDAGQYRCVAKNSLGTAYSKVVK 205
                                                    206 LEV 208
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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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US-08-975-307B-11
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US-08-47-464-3
US-08-374-834-16
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Sequence 1, Appli	Sequence 31, Appl	41,	Sequence 2, Appli	Sequence 46, Appl	-	Sequence 48, Appl	•	21,	-	Sequence 15, Appl	•	•	•	Sequence 57, Appl	•	Sequence 43, Appl	Sequence 42, Appl

ALIGNMENTS

Sequence 331, Application US/09188930A

| Patent No. 6150502
| GENERAL INFORMATION:
| APPLICANT: Watson, James D.
| APPLICANT: Strachan, Lorna APPLICANT: Sleeman, Matthew APPLICANT: Sleeman, Matthew APPLICANT: Murison, James Greg |
| TITLE OF INVENTION: Compositions Isolated From Skin Cells |
| TITLE OF INVENTION: Compositions Isolated From Skin Cells |
| TITLE OF INVENTION: Compositions Isolated From Skin Cells |
| TITLE OF INVENTION: And Methods For Their Use |
| TITLE OF INVENTION NUMBER: US/09/188,930A |
| CURRENT APPLICATION NUMBER: US/09/188,930A |
| SOFTWARE: FastSEQ for Windows Version 3.0 |
| SEQ ID NO 331 |
| LENGTH: 299 | Query Match 25.9
Best Local Similarity 32.8
Matches 101; Conservative RHKSSFVI AAVLVTLILLGILVFGIWFAYSRGHFDRTKKGTSSKKVIYSQPS-----ARSEGEF GGVLVVLAVLALITLGICCAYRRGYFINNKQDGES----YKNPGKPDGVNYIRTDEEGDF 302 RAFSNSSYVLNPTTGELVFDPLSASDTGEYSCEARNGYGTPMTSNAVRMEAVERNVGVIV PRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQ-EMEVYDLNIGGII 246 EVKVKLIVLVPPSKPTVNIPSSATIGNRAVLTCSEQDGSPPSEYTWFKDGIVMPTNPKST 180 EIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRAN KFDQGDTTRLVCYNNKITASYEDRVTFL-PTGITFKSVTREDTGTYTC-MVSEEGGNSYG KKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEID RKLLCL----FILAILLCSLALGSVTVHSSEPEVRIPENNPVKLSCAY--SGFSSPRVEW RLRLCARLPDFFLLLLFRGCLIGAVNLKSSNRTPVVQEFESVELSCIITDSQTSDPRIEW 25.9%; 48; Score 424; DB 4; Pred. No. 2.5e-35; B; Mismatches 137; Length 299; Indels 22; Gaps 240 187 62 67

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; PRIOR APPLICATION NUMBER: 60/052,186
; PRIOR FILING DATE: 1997-07-10
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSEQ for Windows Version 3.; SEQ ID NO 2
; SEQ ID NO 2
LENGTH: 299
TYPE: PRT
ORGANISM: HOMO SAPIENS
US-09-462-270-2
            APPLICANT: Watson, James D.

APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Onrust, Rene
APPLICANT: Onrust, Rene
TITLE OF INVENTION: Compositions Isolated From Sk:
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1011c1
CURRENT APPLICATION NUMBER: US/09/188,930A
CURRENT FILING DATE: 1998-11-09
NUMBER OF SEO ID NOS: 348
SOFTWARE: FastSEQ for Windows Version 3.0
SEO ID:NO 189
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                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               Sequence 189, Application US/09188930A Patent No. 6150502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 25.9%; Score 424; DB 4; Length 299; Best Local Similarity 32.8%; Pred. No. 2.5e-35; Matches 101; Conservative 48; Mismatches 137; Indels
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LENGTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            181 RAFSNSSYVLNPTTGELVFDPLSASDTGEYSCEARNGYGTPMTSNAVRMEAVERNVGVIV
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; TYPE: PRT
; ORGANISM: HOMO SAPIENS
US-09-462-270-4
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APPLICANT: SmithKl
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TITLE OF INVENTION: Human Fil Antigen: A Cell Surface
TITLE OF INVENTION: Receptor Involved in Platelet Ag
FILE REFERENCE: GH-70150US
CURRENT APPLICATION NUMBER: US/09/462,270
CURRENT FILING DATE: 2000-01-05
PRIOR APPLICATION NUMBER: 60/052,186
PRIOR APPLICATION NUMBER: 60/052,186
PRIOR FILING DATE: 1997-07-10
NUMBER OF SEQ ID NOS: 4
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ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              303 RHKSSFVI 310
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                                   89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63 KFDQGDTTRLVCYNNKITASYEDRVTFL-PTGITFKSVTREDTGTYTC-MVSEEGGNSYG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGVLVVLAVLALITLGICCAYRRGYFINNKQDGES----YKNPGKPDGVNYIRTDEEGDF 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQ-EMEVYDLNIGGII 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EVKVKLIVLVPPSKPTVNIPSSATIGNRAVLTCSEQDGSPPSEYTWFKDGIVMPTNPKST 180
                 KKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEID 127
                                                                                                             RLRLCARLPDFFLLLLFRGCLIGAVNLKSSNRTPVVQEFESVELSCIITDSQTSDPRIEW 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAVLVTXILLGILVFGIWFAYSRGHFDRTKKGTSSKKVIYSQPS-----ARSEXEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RAFSNSSYVLNPTTGELVFDPLSASDTGEYSCEARNGYGTPMTSNAVRMEAVERNVGVIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEID 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RKLLCL----FILAILLCSLALGSVTVHSSEPEVRIPENNPVKLSCAY--SGFSSPRVEW
KFDQGDTTRLVCYNNKITASYEDRVTFL-PTGITFKSVTREDTGTYTC-MVSEEGGNSYG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRAN
                                                                         RKLLCL----FILAILLCSLALGSVTVHSSEPEVRIPENNPVKLSCAY--SGFSSPRVEW
                                                                                                                                                                                                                                                                                                                                              FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative 48; Mismatches 139;
                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25.2%;
                                                                                                                                                                       17.6%; Score 288; DB 4; Length 205; 33.7%; Pred. No. 1.2e-21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 412; DB 4;
Pred. No. 4.2e-34;
                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Platelet Aggregation
                                                                                                                                                       92;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 5712369
                                                                                                                                                                                                                                                                                   Query Match
Best Local 9
                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (128 EIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRAN 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: LUTELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Old, Lloyd J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                    191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PS,
OPERATING SYSTEM:
                                                                                                                                                           84
                                                                                                                                                                                            19
                                                                                                                                                                                                                               29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Hanson, No. 5712369man D. REGISTRATION NUMBER: 30,946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/597,495B FILING DATE: 02-Feb-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
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RNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCE-EQEMEVYDLNIG----GII 246
                                                   LVPPSKPECGIEGETIIGNNIQLTCQSKEGSPTPQYSWKRYNILNQEQPLAQPASGQP-- 195
                                                                                   QVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDV-----PLPTDSRANPRF 190
                                                                                                                      YIHGELYKNRVSISNNAE-QSDASITIDQLTMADNGTYECSVSLMSDLEGNTKSRVRLLV
                                                                                                                                                       -IQGDL-----AGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELTV 135
                                                                                                                                                                                            VDAISVETPODVLRASQGKSVTLPCTYHTSTSSREGLIQWDKLLLTHTERVVIWPFSNKN 78
                                                                                                                                                                                                                             IGAVNLKSSNRTPVVQEFESVELSCII-TDSQTSDPRIEWKKIQDEQTTYVF---FDNK- 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RAFSNSSYVLNPTTGELVF 199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ESSEE: Felfe & Lynch
ET: 805 Third Avenue
: New York City
E: New York
                                                                                                                                                                                                                                                                  Similarity. 25.79; Conservative
                                                                                                                                                                                                                                                                                                                                                                                           amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Simpson, Richard J.; Nice, Edouard; Moritz, R. L.; Simpson, Richard J.; Nice, Edouard; Moritz, R. L.; Catinel, B.; Ji, Hong; Burgess, Anthony W.; Heath, Joan K.; White, Sara J.; Johnstone, Camprovention: Colon Cell And Colon 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (212) 838-3884
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (212)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 688-9200
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                                                                                                                                                                                                                                                                14.0%; Score 228.5; DB 1
25.7%; Pred. No. 2.7e-15;
tive 49; Mismatches 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LUD 5316.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inch, 360 kb storage
                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                     319;
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US-09-068-051A-22
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GENERAL INFORMATION:
                                                                                                                   Matches
                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           247
                                                                       29 IGAVNLKSSNRTPVVQEFESVELSCII-TDSQTSDPRIEWKKIQDEQTTYVF---FDNK- 83
     84 -IQGDL-----AGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELTV 135
                                        19 VDAISVETPQDVLRASQGKSVTLPCTYHTSTSSREGLIQWDKLLLTHTERVVIWPFSNKN 78
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/597,495
FILING DATE: 02-Feb-1996
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: PC-
SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10103
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Fulbright & Jaworski LLP
STREET: 666 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Old, Lloyd J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Colon Cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EGDFRHK 305
                                                                                                                                   Similarity
                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212)
                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Hanson, No. 6291235man D. REGISTRATION NUMBER: 30,946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/068,051A FILING DATE: 10-Dec-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette, 3.5 inch, 360 kb storage COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York City
STATE: New York
                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212)
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Old, Lloyd J.; Welt, Sydney; Ritter, Gerd; Simpson, Richard J.; Mice, Edouard; Moritz, Catimel, B. Ji, Hong; Burgess, Anthony W.; Heath, Joan K.; White, Sara J.; Johnstone,
                                                                                                                                                                                                                                                                                   319 amino acids
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                                                                                                                                   14.08;
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                                                                                                                                                                                                                                                                                                                                           752-5958
                                                                                                                                                                                                                             SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PC-DOS
                                                                                                                 49;
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                                                                                                                                   Score 228.5;
Pred. No. 2.
                                                                                                                   Mismatches
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                                                                                                                                   5; DB 4;
2.7e-15;
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                                                                                                                 Indels
                                                                                                                                                  Length 319;
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                                                                                                                 51;
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                                                                                                               Gaps
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136 QVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDV-----PLPTDSRANPRF 190

YIHGELYKNRVSISNNAE-QSDASITIDQLTMADNGTYECSVSLMSDLEGNTKSRVRLLV

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US-09-068-051A-32; Sequence 32, Application US/09068051A fatent No. 6291235
GENERAL INFORMATION:
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                                                                                                                                                                                 ; TOPOLOGY: linear; SEQUENCE DESCRIPTION: US-09-068-051A-32
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                                                                                                  Query Match
Best Local 9
                                                                                 Matches
                                                                                                                                                                                                                                                                                   TELEFAX: (212) 752-51
INFORMATION FOR SEQ ID NO: 32
SEQUENCE CHARACTERISTICS:
                                     48 SVELSCIITDSQTSDPR--IEWKKIQDEQT-----TYVF---FDNKIQGDLAGR 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         196
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                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/597,495
FILING DATE: 02-Feb-1996
APPLICATION NUMBER: 08/511,876
FILING DATE: 04-Aug-1995
ATTORNEY/AGENT INFORMATION:
NAME: Hanson, No. 6291235man D.
REGISTRATION NUMBER: 30,946
REGISTRATION NUMBER: LUD 5316.2
TELECHONUS: (212) 318-3168
TELEFAX: (212) 752-5958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               298
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                                                                               Local Similarity hes 72; Conserv
    37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 360 kb storage
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WOrdperfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/068,051A
FILING DATE: 10-Dec-1998
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSE: Fulbright & Jaworski LLP
STREET: 666 Fifth Avenue
CITY: New York City
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Old, Lloyd J.; Welt, Sydney; Ritter, Gerd;
Simpson, Richard J.; Nice, Edouard; Moritz, R. L.;
Catimel, B.; Ji, Hong; Burgess, Anthony W.;
Heath, Joan K.; White, Sara J.; Johnstone, Cameron
TITLE OF INVENTION: Colon Cell And Colon Cancer Cell
Associated Nucleic Acid Molecules, Protein And Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LVPPSKPECGIEGETIIGNNIQLTCQSKEGSPTPQYSWKRYNILNQEQPLAQPASGQP--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VGVVAALITIGITIYCCCC---RGKDDNTEDKEDARPNREAYEEP--PEQLRELSREREE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCE-EQEMEVYDLNIG---GII 246
SVTLPCTY-NTYVSDREGFIQWDKLLRSQTERVVTWNFVTKKYIYGNRYENRVR--VSND 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EDDYRQE 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EGDFRHK 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -----VSLKNISTDTSGYYICTSSNEEGTQFCNITVAVRSPSMNVALYVGIA 242
                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 10103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                 LENGTH: 318 amino acids
                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                                                                                                    13.2%; Score 215.5; DB 4; 24.1%; Pred. No. 5.8e-14;
                                                                                                                                                                                                        SEQ
                                                                                 40;
                                                                                                                                                                                                        IJ
                                                                                 Mismatches
                                                                                                                                                                                                        NO:
                                                                                                                                                                                                          32
                                                                                                                     Length 318;
                                                                                 Indels
                                                                                 73;
                                                                             Gaps
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RESULT 8
US-09-175-928-2
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APPLICANT: Jacobs, Kenneth
APPLICANT: MCCOy, John M.

APPLICANT: Cavallie, Edward R.

APPLICANT: Cavallie, Edward R.

APPLICANT: Sans, Cheryl
APPLICANT: Treacy, Maurice
APPLICANT: Treacy, Maurice
APPLICANT: Mi, Sha
APPLICANT: Genetics Institute, Inc.
APPLICANT: Genetics 1006B ANJI72A
FILE REFERENCE: 6006B ANJI72A
FILE REFERENCE: 6006B ANJI72A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/175,928A
CURRENT FILING DATE: 1998-10-20
NUMBER OF SEQ ID NOS: 62
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 387
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 12.1%; Score 197.5; DB 4; Best Local Similarity 25.4%; Pred. No. 5.4e-12; Matches 76; Conservative 43; Mismatches 119;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                               68 --KKIQDEQTTYVF-----FDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRC
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                                                                                                                            ---NDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEE 232
                                                                                                                                                                                                                                                                                                                                                                             FWKVFLILSCLAGQVSVV----QVTIPDGFVNVTVGSNVTLICIYTTTVASREQLSIQWSF 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VGKMATLHCQESEGHPRPHYSW-----YRNDVPLPTDSRANPRFRNSSFHLNSETGTLVF 206
                                                                                          LEGRDI-VPVKENFNP---
                                                                                                                                                                                         DVNNPPDFLGQNQGILNVSVLVKPSKPLCSVQGRPETGHTISLSCLSALGTPSPVYYWHK 176
                                                                                                                                                                                                                                                                                FHKKEMEPISIYFSQGGQAVAIGQFKDRITG----SNDPGNASITISHMQPADSGIYIC
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                                      QEMEVYDLNIG----GIIGGVLVVLAVLALITLGICCAYRRGYFINNKQDGESYKNPGK 287
                                                                                                                                                                                                                                  EVVARNDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYR 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IVYCCCC-----REKDDKDQDREDARPNRAAYQVPKKEQKEISRGREDED-DHRHE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IGNNIQLTCHSAEGSPSPQYSWKSYNAQNQQRPLTQPVSGEP--
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ITLGICCAYRRGYFINNKQDGESYKNPGKPDGVNYI-----
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---IDLTSSHPEVGIIVGALIGSLVGAAIIISVVC-----FARNKAKAKAKERNSK 271
                                                                                            ---TTGILVIGNLTNFEQGYYQCTAINRLGNSSCE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 387;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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RESULT 9 US-08-979-424-1

Sequence 1, Application US/08979424 Patent No. 5942606

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RESULT 10
US-08-928-383B-2
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Sequence 2, Application US/08928383B Patent No. 6210921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/979,424
FILING DATE: Filed Herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: VIRAL RECEPTOR PROTEIN
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 390 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                          119
                                                                                                                                              222 SNDAGSARCEEQEMEVYD----LNIGGIIGGVLVVLAVLALITLGICCAYRRG 270
                                                                                                                                                                                                                       165 GHPRPHYSWYRNDVPLPTDSRANPRFRNSSFH---LNSETGTLVFTAVHKDDSGQYYCIA 221
                                                                                                                                                                                                                                                                                             108 RDSALYRCEV-VARNDRKEIDEIV--IELTVQVKPVTPVCRVPKAVPVGKMATLHCQESE 164
                                                                                                                                                                                                                                                                                                                            62 SSQPWEVPFVMWFFKQKEKEDQVLSYI---NGVTTSKPGVSLVYSMPSRNLSLRLEGLQE 118
                                                                                                                                                                                                                                                                                                                                                                     57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LIBRARY: LUNGF
CLONE: 1232054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 650-855-05
TELEFAX: 650-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                        22 ALAPPSRAQLQLHLP--------ANRLQAVEGGEVVLPAWYTLHGEVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Incyte Pharmaceuticals, STREET: 3174 Porter Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                                            2 ALRRPPRLRLCARLPDFFLLLLFRGCLIGAVNLKSSNRTPVVQEFESV-----ELSCIIT 56
                                                                                                                                                                                  SKPAVQYQW-----DRQLPSFQ--TFFAPALDVIRGSLSLTNLSSSMAGVYVCKA
                                                                                                                                                                                                                                                           KDSGPYSCSVNVQDKQGKSRGHSIKTLELNVLVPPAPPSCRLQGVPHVGANVTLSCQSPR 178
                                                                                                                                                                                                                                                                                                                                                                  DSQTSD-PRIEW----KKIQDEQTTYVFFDNKIQGDLAGRAEILG----KTSLKIWNVTR 107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10.4%; Score 169.5;
26.3%; Pred. No. 4e-(
tive 37; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              116;
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                                                                                                              275
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APPLICANT:

Lal, Preeti Corley, Neil C

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RESULT 11
US-08-979-424-3
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Sequence 3, Application US/08979424 Patent No. 5942606 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 9.8%; Score 160.5; DB 4; Best Local Similarity 26.3%; Pred. No. 3e-08; Matches 74; Conservative 33; Mismatches 123;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE: 365 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/026
FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: DFN-
TELECOMMUNICATION INFORMATION:
TELECHNONE: (61),227,7400
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/928,383B
FILING DATE: 12-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein
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APPLICANT: and Marshall S. Horwitz
TITLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
TITLE OF INVENTION: Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                        224 LLRLNVVPPSNKAGLIAGAIIGTLLALALIGLIIFCCRKKR 264
                                                                                                                                                                                               231 EEQEMEVYDLNIGGIIGGVLV-VLAVLALITLGI-CCAYRR 269
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                                                                                                                                                                                                                                                                    WYR--NDVPLPTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARC
                                                                                                                                                                                                                                                                                                                               KV-----KKAPGVANKKIHLVVLVKPSGARCYVDGSEEIGSDFKIKCEPKEGSLPLQYE
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US-09-272-496-2; Sequence 2, Application US/09272496; Patent No. 6245966; GENERAL INFORMATION:
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Best Local Sim
Matches 74;
APPLICANT: DeGregori, TITLE OF INVENTION: A
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NAME: Billings, Lucy J

REGISTRATION NUMBER: 36,749

REFERENCE/DOCKET NUMBER: PF-0

TELECOMMUNICATION INFORMATION:

TELECHONE: 650.855-5555
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LIBRARY: GenBank
CLONE: 1946351
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   68 --KKIQDEQTTYVFFDNKIQG----DLAGRAEIL-----GKTSLKIWNVTRRDSALYRC 115
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TOPOLOGY: 11
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                                                                                                                                                                             RLNVVPPSNKAGLIAGAIIGTLLALALIGLIIFCCRKKR 264
                                                                                                                                                                                                                                                                                                                                                                      EVVARNDRKEIDEIV---IELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYS
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  Adenoviral mediated
                    James
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gene transfer into lymphocytes
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; SEQ ID NO 2
; LEWGTH: 365
; TYPE: PRT
; ORCANISM: Homo sapiens
US-09-272-496-2
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Best Local 9
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CURRENT APPLICATION NUMBER: US/09/272,496
CURRENT FILING DATE: 1999-03-19
EARLIER APPLICATION NUMBER: US 60/092782
EARLIER FILING DATE: 1998-07-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SWEET, RAYMOND
APPLICANT: TRUNGH, ALEMSEGED
TITLE OF INVENTION: A HUMAN I
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                   APPLICATION NUMBER: 60/0:
FILING DATE: 22-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: PRESTIA, PAUL F
                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
REFERENCE/DOCKET NUMBER: GI
                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                              STREET: P.U. C. CITY: VALLEY FORGE
                                     REGISTRATION NUMBER:
                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 08-DEC-1997
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les 74; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RLNVVPPSNKAGLIAGAIIGTLLALALIGLIIFCCRKKR 264
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O. BOX 980
                                                                                                                                                                                                                                                                                       Diskette
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                                                                                                                60/059,448
                                 23,031
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                   GH-70264
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Pred. No. 3.6
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nes 123;
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

2:

LENGTH:

1101 amino acids

TELEPHONE:

610-407-0700

TELEFAX:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 11, Applicat Patent No. 5952171
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: McCarthy, Sean A.
APPLICANT: Gearing, David P.
APPLICANT: Levinson, Douglas
                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    864 VV 865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     810 SSIVLTSLVWVCIIYQ----TRKKSEEYSVTNTDETVVPPDVPSYL--SSQGTLSDRQET 863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                144 CRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPLPTDSRANPRFRNSSFHLNSETGT 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           253 LAVLALITLGICCAYRRGYFINNKQDGESYKNPGK----PDGVNYIRTDEEGDFRHKSSF 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              204 LVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDLNI------GGIIG--GVLVV 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        706 PLEDRVVSVGETVALQC-KATGNPPPRITWFKGDRPLSLTER------HHLTPDNQL 755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             606 LPSFTKTPHDITIRTTTVARLECAATGH---PNPQIAWQKDGGTDFPAARERRMHVMPDD- 662
                                                                                                   APPLICATION NUMBER: US/O FILING DATE: 19-NOV-1996
                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEO for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                          COUNTRY: US
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84 IQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELTVQVKPVTPV 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34 LKSSNRTP---VVQEFESVELSCIITDSQTSDPRIEWKK-----IQDEQTTYVFFDNK 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 70; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LVVQNVVAEDAGRYTCEMSNTLGTERAHSQ-----LSVLPAAGCRKDGTTVGIFTIAVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -----DVFFITDVKI-----DDAGVYSC--TAQNSAGSISANA-TLTVLETPSLVV 705
                                                                                                                                                                                                                                                                                                                                        Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/08752307B
                                                                                                                                                                                                                                                                                                                                                           225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
                                                                                                                                                                                                                                                                                                                                                                                                                                           METHOD FOR IDENTIFYING GENES
ENCODING NOVEL SECRETED OR MEMBRANE-ASSOCIATED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Douglas A.
                                                                                                                      US/08/752,307E
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Best Local S
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             TELEFAX: (619) 554-63
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Crossin, Kathryn L.
TITLE OF INVENTION: NEURITE OUTGROWTH-PROMOTING
TITLE OF INVENTION: CONTAINING FIBRONECTIN TYPE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 094
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
SEQUENCE CHARACTERISTICS:
                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                              CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: V. TELEFAX: 200154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        248 TPMGSTSNKVELRGNVLLLECIAAGLPT--PVIRWIKEGGELPANRTFFEN-----
                                                                              REGISTRATION NUMBER: 34
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                 CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92
                                             TELEPHONE:
                                                                                                                                                              APPLICATION NUMBER: US/0 FILING DATE: 24-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40 TPVVQEFESVE-----LSCIITDSQTSDPRIEWKKIQDE-QTTYVFFDNKIQGDLAGR 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE:
                                                                                                               NAME:
                                                                                                                                                                                                                                                                                                                 COUNTRY:
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les 60; Conserv
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AVQERSSAVYQCNASNEYG 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AVHKDDSGQYYCIASNDAG 226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VGKMATLHCQESEGHPRPHYSWYRNDVPL---PTD-SRANPRFRNSSFHLNSETGTLVFT 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEILGKTSLKIWNVTRRDSALYRCEVVARNDRKEIDEIVIELTVQVKPVTPVCRVPKAVP
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                                                                                                           Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          612 amino acids
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10550 No. 6313265th Torrey Pines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cunningham,
                                                                                                                                                                                                                                                                                                                 U.S.
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                                              (619) 554-2937
                           554-6312
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2: 35,283
                                                                                                                                                                               US/08/506,296B
                                                                                          34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bruce A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21;
                                                                               TSRI
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                                                                               488.0
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                                                                                                                                                                                                                                                                                                                                                                  Road,
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III REPEATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 612;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    AND METHODS
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OF USE

Search completed: September 19, 2002, 17:33:16
Job time: 234 sec

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OM protein - protein search, using sw model
                                                                                             GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Run on: September 19, 2002, 17:40:17; Search time 30.81 Seconds (without alignments) 966.819 Million cell updates/sec

Title: Perfect score: Sequence: US-09-524-531C-15 310 MALRRPPRLRLCARLPDFFL......VNYIRTDEEGDFRHKSSFVI 310

Scoring table: OLIGO Gapop 60.0 , Gapext 60.0

Searched: 283138 seqs, 96089334 residues

size :

0

Total number of hits satisfying chosen parameters: 283138

Minimum DB Maximum DB seq length: 0 seq length: 2000000000

Post-processing: Listing first 75 summaries

Database :

PIR_71:*

pir1:*
pir2:*
pir3:*
pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25 26 27 28 29	2222	16 17 18 19	13	11 9 11	54321	Result No.
7777	17777	17777	7777	7 & & & & & & & & & & & & & & & & & & &	~~~~	Score
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0		, , , , , , , , , , , , , , , , , , ,		~ @ @ @ @ @ ^ % % % % % %	00000 00000	Query Match
242 244 255 280 294	208 211 214 217	173 173 177 177 177 208	113 171 172 173	366 366 492 784	290 336 342 366	Length
88888	00000	88888	00000		22422	B
G84315 A46066 T35883 PH0269 H84023	T01588 E86888 T47268	B90241 T25730 E85833 B90988 T03627	AH1748 G70548 T08548 T01282	JH0547 JH0545 I37078 AB1013 A86676	I68750 A86406 HLHUC4 E82933 JH0546	Ib
cobalt transport p lymphotoxin beta - transcription regu epidermal autoanti phosphate ABC tran	B	p ba	hypothetical prote hypothetical prote hypothetical prote hypothetical prote hypothetical prote	class I histocompa class I histocompa class I histocompa HLA-C alpha chain probable membrane carbon starvation	MHC class I lympho probable RING zinc MHC class I histocompa class I histocompa	

probable Abc class	2 B87098		3 558	٠.	٦,	4. IU
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tyrosine-specific				٠	7	Ģ
tyrosine-spec	G909			٠	7	4
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endosperm specific	T0434				7	-
hypothetical					7	0
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hypothetical prote					7	7
hypothetical	T2496				7	σ
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hypothetical prote	F7181				7	4
hypothetical	T4637			•	7	ω
GGDEF family	G7561			•	7	N
a	D7530			•	7	_
robable s	н9587			•	7	0
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hypothetical	T2730				7	æ
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sugar transport	C9819				7	0
lymphotoxin-beta	14913	σ	w		7	G
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tra	A9824	9	2		7	N
nypothetical	ABSUS					-

ALIGNMENTS

RESULT 1
168750
MHC class I lymphocyte antigen - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 07-Jun-1996 #sequence_revision 07-Jun-1996 #text_change 21-Jan-2000

C;Accession: 168750

R;Pohla, H.; Kuon, W.; Tabaczewski, P.; Doerner, C.; Weiss, E.H.

Immunogenetics 29, 297-307, 1989

A;Title: Allelic variation in HLA-B and HLA-C sequences and the evolution of the HLA-A;Reference number: 154457; MUID:89233295

A;Accession: 168750

A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-290 <RES>
A;Cross-references: GB:M28207; NID:g576478; PIDN:AAA53259.1; PID:g576479
C;Superfamily: class I histocompatibility antigen; immunoglobulin homology
F;144-209/Domain: immunoglobulin homology <IMM>

Query Match 2.6%; Score 8; DB 2; Best Local Similarity 100.0%; Pred. No. 5.8; Matches 8; Conservative 0; Mismatches Length 290

0; Indels 0; Gaps

0

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C;Accession: A86406
R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alons C, Thin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, F. Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, F. Chasen, N.F.; Hughes, B.; Hulzar, L. Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marzial C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marzial Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.; Southwick, A.M.; Sun, H.; Tallor M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A;Ritle: Sequence and analysis of chromosome 1 of the plant Arabidopsis.

A;Reference number: A86141; MUID: 21016719
                        밁
                                                                                                                                                                                                                                A;Gene: GDB:HLA-C
A;Cross references: GDB:119311; OMIM:142840
A;Map position: 6p21.3-6p21.3
C;Superfamily: class I histocompatibility antigen; immunoglobulin | C;Keywords: glycoprotein; heterodimer; transmembrane protein; trans E;196-261,Domain: immunoglobulin homology <IMM>
E;186/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references:
C;Genetics:
A;Map position: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probable RING zinc finger protein [imported] - Arabidopsis thaliana
C; Species: Arabidopsis thaliana (mouse-ear cress)
C; Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 31-Mar-2001
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C;Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-342 < DAV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Homo sapiens (man)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-336 <STO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: A24512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MHC class I histocompatibility antigen HLA-C4 alpha chain -
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                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   iol. Chem. 260, 13414-13423, 1985
Ttle: Comparison of HLA class I gene sequences.
eference number: A92500; MUID:86033791
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                                                                        249 VLVVLAVL 256
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                        VLVVLAVL 300
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                                                                                                                                                                                                                                                                                                                                                                                                                        GB:M11886; NID:g184173; PIDN:AAA52665.1; PID:g386777
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100.0%;
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                                                                                                                                       Score 8; Pred. No.
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Pred. No.
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5. 6.7;
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6 N
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ne protein; transplantation
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war, K.
class I histocompatibility antigen Gogo-C0203 heavy chain precursor - lowland
C;Species: Gorilla gorilla gorilla (lowland gorilla)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 23-Jul-1999
                                                                                                                                                               В
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A; Introns: 25/1; 115/1; 207/1; 299/1; 339/1; 350/1; 36(Superfamily: class I histocompatibility antigen; immu C; Keywords: transmembrane protein F;1-24/Domain: signal sequence #status predicted <SIG>F;25-366/Product: class I histocompatibility antigen hu F;25-114/Domain: alpha-1 <ALI>F;25-114/Domain: alpha-1 <ALI
F;25-114/Domain: alph
                                                                                                                                                       Query Match
Best Local Similarity
Watches 8; Conserv:
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F;207-298/pomain: alpha-3 <AL3>
F;220-285/pomain: immunoglobulin homology
F;220-285/Domain: intracellular #status pr
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A;Title: Gorilla class I major histocompatibility A;Reference number: JH0534; MUID:92078860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                class I histocompatibility antigen Gogo-C0202 heavy chain precursor - C;Species: Gorilla gorilla gorilla (lowland gorilla) C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 23-Jul C;Accession: JH0546
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A; Genetic code:
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A;Experimental source: serovar 3; blovar 1
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A; Residues: 1-348 <GLA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 18-Aug-2000 #sequence_revision
C;Accession: E82933
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Best Local :
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324
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Pred. No.
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O92114 mus musculu
O06643 homo sapien
C67798 aguifex aeo
P41155 mus musculu
O48146 homo sapien
O50840 methanococc
P18199 escherichia
P32506 cercopithec
P15151 homo sapien
P80595 solanum tub
P46104 streptomyce
P00178 oryctolagus
P11168 homo sapien
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. MEDLINE=86033791; PubMed=3863816;
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Pfam; PF00129; MHC_I; 1.
PfoDom; PD000050; MHC_I;
SMART: SM00407; IGC1; 1.
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J. Exp. Med. 174:1491-1509(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gorilla gorilla gorilla (Lowland gorilla)
Eukaryota; Metazoa; Chordata; Craniata; V
Mammalia; Eutheria; Primates; Catarrhini;
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01-NOV-1997 (Rel. 35, Last annotation update)
Class I histocompatibility antiqen, GOGO-C0201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          use by non-profit institutions as long modified and this statement is not removed entities requires a license agreement (See
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InterPro; IPR003597;
InterPro; IPR001039;
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PIR; JH0545; JH0545.
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CLASS I HISTOCOMPATIBILITY ANTIGEN,
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Q9my31 homo sapien
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Q9c7e9 arabidopsis
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InterPro; IPR003006; Ig_MHC.
Pfam: PF00047; 1g; 2.
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SMART; SM00409; IG; 2.
SMART; SM00401; IG; 1.
Immunoglobulin domain; Signal.
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SEQUENCE FROM N.A., Johnson-leger C., Wong C., DuPasquie
Aurrand-Lions M.A., Johnson-leger C., Wong C., DuPasquie
"Heterogeneity of endothelial junctions is reflected by
""""" and specific subcellular localization of the
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"Junctional adhesion
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Metazoa; Chordata;
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Mammalia; Eutheria;
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Matches 308
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Mammalia; Eutheria; Ro
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Catarrhini; Hominidae;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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inflammation; cancer; wound; angiogenesis; human;
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10-MAR-1999;
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   GENENTECH
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99WO-US21090

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99US-0162506

99WO-US28214

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99WO-US28301

99WO-US28551

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99WO-US28565
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99US-0146222.
99WO-US20111.
99WO-US20594.
99WO-US20944.
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99US-0141037.
99US-0144758.
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99US-0128849.
99WO-US08615.
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99US-0132371
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RESULT
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DT 26-S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes sixty four human PRO proteins which can CC be used in the treatment of immune related diseases. The human PRO CC proteins, anti-PRO antibodies, agonists and antagonists are useful for CC treating and diagnosing immune related disorders. The disorders are CC selected from systemic lupus erythematosus, rheumatoid arthritis, somethic selected from systemic lupus erythematosus, rheumatoid arthritis, systemic selected; and channed the chronic arthritis, systemic systemic vasculitis, sarcoidosis, autoimmune haemolytic CC systemic selected; and commune thrombocytopaenia, thyroiditis, diabetes mellitus, CC immune-mediated renal disease, demyelinating diseases of the central CC and peripheral nervous systems, hepatobiliary diseases, inflammatory CC bowel disease, gluten-sensitive enteropathy and Whipple's diseases, CC autoimmune or immune-mediated skin diseases, allergic diseases, CC autoimmune or immune-mediated skin diseases, allergic diseases, CC immunological diseases of the lung, and transplantation associated CC diseases including graft rejection and graft-versus-host-disease. CC AAC58397 to AAC58578 represent PCR primers and hybridisation probes used in the isolation of human PRO sequences. AAC5859 to AAC58642 and CC AAB33414 to AAB3477 represent human PRO polynucleotide and protein cross sequences given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                     AAY96735;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sixty four PRO polypeptides, useful in the diagnosis and treatment of immune related disorders, e.g. systemic lupus erythematosis, rheumatoid arthritis, osteoarthritis, thyroiditis and diabetes mellitus -
   26-SEP-2000
                                                                      AAY96735 standard; Protein; 310
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Kabakoff RC,
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Lu Y,
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Pan J, Pennica
D, Watanabe CK,
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Pred. No. 1.7e-295;
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D, Shelton DL,
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DL, Smith
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12-JAN-1999
12-JAN-1999
12-JAN-1999
12-JAN-1999
12-FEB-1999
10-FEB-1999
10-FEB-1999
12-FEB-1999
                                                                                                                       Botstein D,
Gurney AL,
This is PRO1868, a putative homologue of A33 antigen, a known colorectal cancer-associated marker. The invention concerns novel secreted and transmembrane proteins, designated PRO polypeptides. The
                                       Claim 1;
                                                       Isolated nucleic acid molecule encodes transmembrane polypeptide
                                                                                   WPI; 2000-431586/37.
N-PSDB; AAA51265.
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anti-inflammatory;
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                                      Fig 14; 154pp; English.
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an J, Roy MA,
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9805-0113145
9805-0113511
9805-0113511
9905-011558
9905-0115733
9905-0119341
9905-0119955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Tyrosine kinase phosphorylation site"
104..107
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cytostatic;
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                                                                                                                                 Ferrara N,
                                                                                                                        Stewart TA,
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                                                                                                                       Fong S,
A, Tumas
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D, Watanabe
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RESULT
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Best Local Sim
Matches 310;
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11-JAN-2000;
02-MAY-2000;
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                                                                                                                                  0ta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-NOV-2001
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  830 Primers useful for
                                           N-PSDB;
                                                             WPI; 2001-524255/58
                                                                                                                                                                                                                                                                                                            07-JUL-2000; 2000EP-0114089
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                                                                                                                                  Nishikawa T,
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; 2000JP-0118774.
; 2000JP-0183765.
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na T, Nagai
  synthesizing
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Pred. No. 1.7e-295;
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K, Kojima
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S, Otsuki
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11-JAN-2000;
02-MAY-2000;
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                                                                                                                 07-JUL-2000;
                                                                                                                                                                                                                                                                                                        Human;
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                                                                                                                                                                                                                                                                                                                                                   polypeptide,
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Pred. No. 1.7e-295;
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RESULT
AAU12440
ID AAU1
XX
AC AAU1
XC AAU1
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DT 24-O
XX
Huma
XX
Huma
KW Huma
KW brea
KW cart
KW adip
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                        breast; prostate; cervical; tumour cartilage; ear; proliferation; gluadipocyte; A-peptide; factor VIIA;
                                                                                                                                                                                                              24-OCT-2001
                                                                                                                                                                                                                                                                                                                     AAU12440 standard; Protein;
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310; Conserv
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                                                                                                                                                          polypeptide sequence
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                        nd transmembrane; PRO; mammalian; cancer; lung; cervical; tumour necrosis factor-alpha; TNF-alpha; roliferation; glucose; free fatty acid; skeletal muscle; ide; factor VIIA; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 310; DB 22;
100.0%; Pred. No. 1.7e-295;
tive 0; Mismatches 0;
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1 T, Nagai K, Kojima S,
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S, Otsuki T, Koga
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24-FEB-2000;
01-MAR-2000;
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21-MAR-2000;
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17-MAY-2000;
22-MAY-2000;
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30-DEC-1999
11-FEB-2000
11-FEB-2000
         of T-lymphocytes, the release of a cytokine from peripheral blood monocytes (PBMCs), or the proliferation of endothelial cells. Some of the PRO polypeptides may modulate glucose or free fatty acid uptake by skeletal muscle cells or by adipocytes; or inhibit binding of A-peptide to factor VIIA. The PRO polypeptides can be used in assays to identify molecules involved in binding interactions. The polynucleotides encoding PRO polypeptides can be used to generate probes, antisense RNA/DNA, transgenic or knock out animals and can be used in gene therapy.
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Smith V,
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18-FEB-2000;
22-FEB-2000;
                                                                                                                                                               PRO polypeptides, and to detect the presence of mammallan lung, colon, breast, prostate, rectal, cervical or liver tumours by comparing PRO polypeptide expression in a cell sample to that in a control sample. Some of the 275 sequences are also useful to stimulate the release of tumour necrosis factor-alpha (TNF-alpha) from human blood, the proliferation or differentiation of chondrocytes, the proliferation or
                                                                                                                                  gene expression in pericyte cells, the release of proteoglycans from cartilage, the proliferation of inner ear utricular supporting cells or
                                                                                                                                                                                                                                                            PRO
                                                                                                                                                                                                                                                                                                                                                                                  Isolated, secretory and transmembrane PRO polypeptide used to detect other PRO polypeptides, link bioactive molecules to cells expressing PRO polypeptides, and detect the presence of mammalian tumours e.g.
                                                                                                                                                                                                                                                                               PRO
                                                                                                                                                                                                                                                                                           PRO
                                                                                                                                                                                                                                                                                                                                        Claim 12; Fig 538; 813pp; English.
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02-JUN-2000;
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                                                                                                                                                                                                                                                          12172-AAU12446 represent novel human secretory and transmembrane polypeptides. The PRO polypeptides are useful to detect other polypeptides, to link bioactive molecules to cells expressing polypeptides, to modulate biological activities of cells expressing
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DB; AAS21512.
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2000WO-US00277
2000WO-US003565
2000WO-US04341
2000WO-US04342
2000WO-US04414
2000WO-US04414
2000WO-US05601
2000WO-US05601
2000WO-US05601
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99US-0170262.
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99WO-US30911
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Tumas D,
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A, Godowski PJ, Gurney
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08-SEP-1999
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05-OCT-1999
05-OCT-1999
29-NOV-1999
16-DEC-1999
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20-DEC-1999
                                                                                                                                                                                                                                                        Human; PRO; dermatological; antipsoriatic; cytostatic; antiinflammatory; antiparkinsonian nootropic; neuroprotective; vulnerary; cardiant; antiangiogenic; vasotropic; antiasthmatic; antirheumatic; cancer; antiarthritic; antiinfertility; antidiabetic; antiviral; diabetes; ophthalmological; gene therapy; skin disease; gastrointestinal disorder;
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26-JUL-1999;
                                                                                                                                                                                                                                                                                                                   Human PRO1868 protein
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(GETH
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                99WO-US23089.
99WO-US28214.
99WO-US28313.
99WO-US30095.
99WO-US30911.
99WO-US30999.
99WO-US00219.
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                                                                                                          99WO-US20594.
99WO-US20944.
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99US-0145698.
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99WO-US21547.
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Pred. No. 1.7e-295;
Mismatches 0;
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RESULT AAB80383

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AAB80383 standard; protein;

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Secreted 24-APR-2001 AAB80383;

encoded human;

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protein; protein

autoimmune; by gene

hyperproliferation;

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung squamous cell carcinoma) and neurodegenerative diseases (e.g. Alzheimer's disease).
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Filvaroff E, Fong S, Gao
Godowski PJ, Grimaldi CJ,
Mather JP, Pan J, Paoni N
                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                            2001-081051/09.
DB; AAF72433.
           DFRHKSSFVI
                                                                                                         PTDSRANPRFRNSSFHLNSETGTLVFTAVHKDDSGQYYCIASNDAGSARCEEQEMEVYDL
                                                                                                                                                        NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL
dfrhkssfvi
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o W, Gerber H,
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NF, Roy MA, S
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Pred. No. 1.7e-295;
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Stewart TA,
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Tumas D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases and microorganism infections
           AAB80408
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                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                   DFRHKSSFVI 310
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                              standard;
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                             protein;
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Olsen HS,
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Pred. No. 1.7e-295;
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Komatsoulis GA;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders
                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                       caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 11; Page 557-558; 601pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200107459-A1
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dfrhkssfvi
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                                          niggiiggvlvvlavlalitlgiccayrrgyfinnkqdgesyknpgkpdgvnyirtdeeg
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                                                                                                                                                                                                                                                                                                                   310 AA;
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                                                                                                                                                                                                                                                                       100.0%;
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Olsen HS,
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                                                                                                                                                                                                                                                                        Score 310; DB 22;
Pred. No. 1.7e-295;
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Birse CE,
                                                                                                                                                                                                                                                              Mismatches
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infection; food.
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Komatsoulis GA;
                                                                                                                                                                                                                                                                                    310;
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                                                                                                                                                                                                                                                                                                                                      nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g corneal infection. Also used in food preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders
                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                  NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL
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cerebrovascular; infection; food.
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Olsen HS,
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Pred. No. 1.7e-295;
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Komatsoulis
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Matches Query Match Best Local

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                                                                                Sequences AAB38321-B38396 represent the amino acid sequences of 62 CC human secreted proteins encoded by the genes AAC69512-C69587. The genes CC and proteins are useful for preventing, ameliorating or treating medical CC conditions, e.g. by protein or gene therapy. The genes are isolated from CC a range of human tissues disclosed in the specification. The nucleic CC acids, proteins, antibodies and (ant)agonists are useful in the CC diagnosis, treatment and prevention of: (a) autoimmune diseases e.g. CC rheumatoid arthritis; (b) hyperproliferative disorders e.g. cardiac CC of the breast or liver; (c) cardiovascular disorders e.g. cardiac CC arrest; (d) cerebrovascular disorders e.g. cardiac CC arrest; (f) nervous system disorders e.g. cardiac CC argiogenesis; (f) nervous system disorders e.g. Alzheimer's disease; (g) infections caused by bacteria, viruses and fungi; and (h) ocular CC disorders e.g. corneal infection. The polypeptides can also be used to CC aid wound healing and epithelial cell proliferation, to prevent skin CC supporting cell culture of primary tissues, to regenerate tissues and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ruben SM, N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 11; Page 603-604; 716pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid molecules encoding 62 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions used as food additives or preservatives -
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26-APR-1999;
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99US-0130991.
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n HS, Ebner
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Sequences AAB38321-B38396 represent the amino acid sequences of 62 human secreted proteins encoded by the genes AAC69512-C69587. The genes and proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the disposis, treatment and prevention of: (a) autoimmune diseases e.g. rheumatoid arthritis; (b) hyperproliferative disorders e.g. neoplasms of the breast or liver; (c) cardiovascular disorders e.g. cardiac arrest; (d) cerebrovascular disorders e.g. cerebra; (e) anglogenesis; (f) nervous system disorders e.g. Alzheimer's disease; (g) infections caused by bacteria, viruses and fungi; and (h) ocular
                                                                                                                                                                         Claim
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                                                                                                                                                                                              Secreted protein; human; autoimmune; hyperproliferation; cardiovascular; cerebrovascular; infection; food.
                                                                                                                                                                                                                                                                                             Gene
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                                                                                                                                                                                                                                                                                                                                                              24-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                              AAB80431;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB80431 standard;
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hes 310;
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                                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                                                                                                                                                                              entry)
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Pred. No. 1.7e-295;
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Best Local S
Matches 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention is used to prevent autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases and microorganism infections •
                                                                                       AAY96294;
                                                                                                          AAY96294 standard;
                    Human;
                                                                16-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HUMA-) HUMAN GENOME
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                                                                                                                                                                                                                                                                                                                                                                                                                                      310;
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Lafleur DW,
                                          IGFAM-6
        immunoglobulin; IGFAM-6; IGFAM; immune disorder;
ion; inflammation; haematopoiesis; AIDS; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                     h 100.0%;
Similarity 100.0%;
10; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 75;
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                                                              (first entry)
                                         immunoglobulin
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Pred. No. 1.8e-295;
; Mismatches 0;
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Birse CE,
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Komatsoulis GA;
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                      cancer;
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Best Local S
Matches 309
                                                                                                                                                                                                                                                                                                                                                                systemic lupus erythematosus and ulcerative colitis, scleroderma, cancer, haemodialysis and extracorporeal circulation, trauma and haematopoietic cancer (such as leukaemia) and infections caused bacteria, viruses, fungi or parasites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IGFAM-6. Its gene was isolated from a cDNA library of leg tissue. It is expressed in reproductive, nervous and cardiovascular tissue, where cancer and inflammation are common. The gene, protein, its antibodies, agonists and antagonists are suitable for diagnosing and treating many diseases, including cancer, immune system disorders (such as inflammation, AIDS, allergies, anaemia, arteriosclerosis, asthma, atherosclerosis, cholecystitis, Crohn's disease, diabetes mellitus, emphysema, Graves' disease, hepatitis, thousands, arteriosclerosis, consistent of the protection of the protect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-NOV-1998;
22-DEC-1998;
07-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunoglobulin superfamily proteins, the agonist and antagonist of protein is useful for preventing and treating disorders associated altered levels of the protein such as cancer, immune system disorder
                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INCY-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Key
  121
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                                                                                                                                                     present sequence is the human immunoglobulin superfamily protein
M-6. Its gene was isolated from a cDNA library of leg
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              NDRKEIDEIVIELTVQVKPVTPVCRVPKAVPVGKMATLHCQESEGHPRPHYSWYRNDVPL
                                                                         SDPRIEWKKIQDEQTTYVFFDNKIQGDLAGRAEILGKTSLKIWNVTRRDSALYRCEVVAR 120
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ndrkeideivieltvqvkpvtpvcrvpkavpvgkmatlhcqeseghprphyswyrndvpl
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99US-0128194.
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238..260
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1..30
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153..221
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Hillman JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hillman
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99.7%;
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                                                                                                                                                                                                                                         Score 209; DB 21;
Pred. No. 1.8e-196;
0; Mismatches 1;
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           antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; and opthalmalogical. The human secreted proteins, polynucleotides, antagonists and agonists of the invention may be useful in the treatment, prevention, and/or diagnosis of various disease, disorders and conditions such as autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, viruses and fungi and ocular disorders e.g. infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to
                                                                                                                                                                                               The polynucleotide sequences given in AAC74223-C74279 encode the human secreted proteins represented in AAB39179-B39226. Sequences AAB39227-B39308 are alternative proteins encoded by the genes, and also protein sequences with which they share homology. The proteins have activities based on the tissues and cells in which they are expressed. Examples of activities include: immunosuppressive; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      opthalmalogical; autoimmune disease; rheumatoid arthritis; angiogenesis; hyperproliferative disorder; cardiovascular disorder; infection; cerebrovascular disorder; nervous system disorder; ocular disorder;
                                                                                                                                                                                                                                                                                                              Disclosure; Page 32;
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-579483/54.
N-PSDB; AAC74237.
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10-DEC-1999;
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antibacterial; virucide; fungicide; neoplasm;
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Best Loc
Matches
The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating
                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity
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23-AUG-2000;
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storage capabilities.
during the isolation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC
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2000US-0649167.
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medical imaging; diagnostic;
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Best Local :
                                                                                             diagnostics, forensite, responsible for genetic
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Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromo and gene mapping, and in recombinant production of (II). The
                                                                                                             New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
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23-AUG-2000; 2000US-0649167.
                                                                       Claim 20;
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upplement; medical imaging; diagnostic;
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                                                                    No 42468; 103pp; English
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Pred. No.
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(II). The

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       uery Match
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New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity
                                                                                                                                                                                                                                              30-MAR-2001; 2001WO-US08631
                                                                                                                                                                                                                                                                                                                                                                          Human; chromosome mapping; gene mapping; gene therapy; forensic
food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human diagnostic protein #22392.
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23-AUG-2000; 2000US-0649167
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                                                                               2001-639362/73.
DB; AAS86588.
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
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Best Local
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WPI; 1999-153781/13
                                  Duclert A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
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                                    Dumas Milne Edwards
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                                                                                                                                                                                                                                                                                                               anti-inflammatory; tumour
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Pred. No.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acids encoding human secreted - proteins obtained from cDNA libraries prepared from substantia nigra, cerebellum, surrenals and fetal brain tissue
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                                                     Drmanac
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          WPI; 2001-639362/73.
N-PSDB; AAS91225.
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                                                                                   (HYSE-) HYSEQ INC.
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23-AUG-2000;
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food supplement; medical imaging; diagnostic; genetic disorder.
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                                                                                                                                                                                                                                                                                                                          diagnostic
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                                                                                                                2000US-0540217.
2000US-0649167.
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Pred. No.
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31-MAR-2000; 23-AUG-2000;

(HYSE-) HYSEQ INC

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2001WO-US08631 2000US-0540217 2000US-0649167

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RESULT
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Best Local
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Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                  .
157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy technique to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                 food si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 20; SEQ ID No 57397; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to isolated polynucleotide (I) and
                                                                                                                                                                  11-OCT-2001
                                                                                                                                                                                                   WO200175067-A2
                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                  Novel human diagnostic protein #7148.
                                                                                                                                                                                                                                                                                                                                                   13-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                   ABG07157 standard; Protein; 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     239 DLNIGGIIGGVLVVLAVLALITLGICCAYRRGYFINNKQDGESYKNPGKPDGVNYIRTDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   299 EGD 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130 dlniggiiggvlvvlavlalitlgiccayrrgyfinnkqdgesyknpgkpdgvnyirtde 189
                                                                                                                                                                                                                                                                 supplement;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  egd
                                                                                                                                                                                                                                                                                 chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   267 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences.
                                                                                                                                                                                                                                                                   medical
                                                                                                                                                                                                                                                                                 mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20.3%;
                                                                                                                                                                                                                                                                 imaging;
                                                                                                                                                                                                                                                                 gene mapping; gene therapy;
maging; diagnostic; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 63; DB Pred. No. 2.3 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 22;
2.3e-53;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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ABG22399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC polypeptide (II) sequences. (I) is useful as hybridisation probes, CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome CC and gene mapping, and in recombinant production of (II). The CC quantifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving CC (II). (II) is useful for generating antibodies against it, detecting or CC quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful in medical disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations in CC diagnostics, forensics, gene mapping, identification of mutations in CC considered the constant protein expression of mutations in CC diagnostics are sequences. ABG00010-ABG30377 represent novel human CC diagnostic amino acid sequences of their brantion.

CC Note: The sequence data for this patent did not appear in the printed constant on the printed specification, but was obtained in electronic format directly from WIPO at figure of the constant of the printed specification, but was obtained in electronic format directly from WIPO at figure and produces.
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WPI; 2001-639362/73
                                   Drmanac RT,
                                                                        (HYSE-) HYSEQ INC.
                                                                                                           31-MAR-2000; 2000US-0540217
23-AUG-2000; 2000US-0649167
                                                                                                                                                                  30-MAR-2001; 2001WO-US08631
                                                                                                                                                                                                         11-OCT-2001
                                                                                                                                                                                                                                           WO200175067-A2
                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                   Human; chromosome mapping; gene mapping; gene therapy; forensic;
food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                          Novel human diagnostic protein #22390.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABG22399 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-639362/73.
N-PSDB; AAS71344.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      214 gdlagraeilgktslkiwnvtrrdsalyrcevvarndrkeideivieltvq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86 GDLAGRAEILGKTSLKIWNYTRRDSALYRCEVVARNDRKEIDEIVIELTVQ 136
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                                   Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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  (I) is useful as hybridisation probes
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human alternation acid sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags
              WPI; 2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity -
                                               Drmanac RT, Liu C,
                                                                                                               31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                  30-MAR-2001; 2001WO-US08631.
                                                                                                                                                                                                                                    WO200175067-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                             (HYSE-) HYSEQ
                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                   Human; chromosome mapping; gene mapping; gene therapy; forension food supplement; medical imaging; diagnostic; genetic disorder
                                                                                                                                                                                                                                                                                                                                                     Novel human diagnostic protein #22389.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     301 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                               2000US-0540217.
2000US-0649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.5%; Score 51;
100.0%; Pred. No.
                                                 YT;
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N-PSDB; AAS86585

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RESULT
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Best Local
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                                                                                                                             Imhof BA,
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Isolated human Confluency Regulated Adhesion Molecule
                                                                WPI; 2000-587436/55
                                                                                                                                                                                                                                                        11-MAR-1999;
                                                                                                                                                                                                                                                                                                                  13-MAR-2000; 2000WO-EP02219
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                                                                                                                                                                                           (RMFD-) RMF DICTAGENE
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inflammation; cancer; wound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human confluency regulated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    family; Ig Sf; vascular adhesion molecule;
; wound; angiogenesis; human;
adhesion molecule 1; CRAM-1; JAM-2.
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Pred. No.
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2.3e-29;
hes 0;
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RESULT 1
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Best Local Similarity
Matches 24; Conserv
                                      The present sequence is the murine confluency regulated adhesion molecu. I (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein and coding sequence can be used in the treatment of cancer, inflammation to modulate cell-cell interactions and angiogenesis, and in the
                                                                                                                                                          Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 or CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                                                                                                                                                                                      WPI; 2000-587436/55.
N-PSDB; AAA97189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin superfamily; Ig Sf; vascular adhesion molecule; inflammation; cancer; wound; angiogenesis; mouse; confluency regulated adhesion molecule 1; CRAM-1; JAM-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB27278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and coding sequence can be used in the treatment of cancer, inflammation, to modulate cell-cell interactions and angiogenesis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                           modulation
                                                                                                                                                                                                                                                                                                                           11-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                     14-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   proteins of the immunoglobulin
                                                                                                                              Example; Fig 8; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                        13-MAR-2000; 2000WO-EP02219
                                                                                                                                                                                                                                                                                                                                                                                                                  WO200053749-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Murine confluency regulated adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in the modulation of wound healing.
                                                                                                                                                                                                                                                                  Imhof BA,
                                                                                                                                                                                                                                                                                             (RMFD-) RMF DICTAGENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       167 PRPHYSWYRNDVPLPTDSRANPRF 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  he present sequence is the human confluency regulated adhesion molecule (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion roteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein nd coding sequence can be used in the treatment of cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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                           of wound
310
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                           healing
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Pred. No.
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4.5e-15;
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Query Match Best Local Similarity

7.7%; 100.0%;

Score 24; Pred. No.

4. BB)B 21; 1.5e-15;

Length

Sequence

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ABB29193
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                                                                       derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genemic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide encoded by a single exon nucleic acid by the method is suitable for the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                             The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic acids
                                                                                                                                                                                                                                                                                                                                                                     New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -
                           Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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27-SEP-2000;
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Sequence
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2000US-0608408.
2000US-0632366.
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2000US-0236359
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Query Match Best Local Similarity

2.6%;

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ABB19769
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                                                                                                                                                                                                                           The invention relates to a single exon nucleic acid probe for measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a peptide encoded by a single exonucleic acid probe of the invention.
                                                                                                                                                                                                                                                                                                                                                             WPI;
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26-MAY-2000;
 ABB19769;
                      ABB19769
                                                                                                                                                                         Sequence
                                                                                                                                                                                           Note: The sequence data for this patent did not form printed specification, but was obtained in electronic from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                        Claim 27; SEQ ID NO 26985; 639pp + sequence listing;
                                                                                                                                                                                                                                                                                                                                         Human
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                             genome-derived single exon nucleic zing gene expression in human fetal
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8; Conserv
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                      standard; Protein;
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2000US-0207456.
2000US-0608408.
2000US-0632368.
2000US-0632368.
2000US-0234687.
2000US-0236359.
2000US-0236359.
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100.0%; Pr
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in electronic form
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liver -
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26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; gene expression; heart; microarray; vascular system; cardiovascular disease; hypertension; cardiac arrhythmia; congenital heart disease.
micmoarray;
                                            05-NOV-2001
                                                                                                                                                                                                                     By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and congenital heart disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                 The present invention relates to single exon nucleic acid probes for measuring human gene expression in a sample derived from human heart (see ABA21335-ABA41305). The present sequence is a protein encoded by one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays.
Humaw; brain expressed exon; gene expression analysis; probe; micxoarray; Alzheimer's disease; multiple sclerosis; schizopi
                            Human brain
                                                             AAM55147;
                                                                                                                                                                                                                                                                                                                                                             Single exon nucleic acid
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                                                                              AAM55147 standard;
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                                                                                                                                          249 VLVVLAVL 256
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                          expressed single exon probe encoded protein SEQ
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2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
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30-JUN-2000;
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                                                                                                                                  Homo
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26-MAY-2000;
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Patent No. 5981730
GENERAL INFORMATION:
APPLICANT: Ausubel, Frederick M.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
APPLICANT: Dahlbeck, Douglas
APPLICANT: Barent, Funiaki
APPLICANT: Kunkel, Barbara N.
APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Windrinos, Michael N.
APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE FAMILY, PR
TITLE OF INVENTION: METHODS
NUMBER OF SEQUENCES: 208
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1
US-08-310-912A-190
  NUMBER OF SEQUENCES: 208
CORRESPONDENCE ADDRESS:
ADDRESSEE: F15h & Rich
STREET: 225 Franklin &
CITY: Boston
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                       Richardson
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APPLICANT: Kunkel, Barbara N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: Mindrinos, Michael N.
APPLICANT: My, Guo-Liang
TITLE OF INVENTION: RESZ GENE FAMILY, PRIMERS, PROBES, AND
TITLE OF INVENTION: DETECTION METHODS
FILE REFERENCE: 00786/254002
CURRENT APPLICATION NUMBER: US/09/301,085
CURRENT FILING DATE: 1999-04-28
EARLIER FILING DATE: 1999-04-28
EARLIER FILING DATE: 1994-09-22
EARLIER APPLICATION NUMBER: 08/227,360
EARLIER APPLICATION NUMBER: 08/227,360
EARLIER APPLICATION NUMBER: 09/227,360
EARLIER FILING DATE: 1994-04-13
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Best Local Similarity
Thes 7; Conserve
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Query Match,
                                                                                                                                               SEQ ID NO 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Ausubel, Frederick M.
                                                                                                                                                                     NUMBER OF SEQ ID NOS: 208
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                       LENGTH: 14
TYPE: PRT
                                                              ORGANISM: Arabidopsis thaliana
)9-301-085-190
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Brent, Andrew F. PPLICANT: Dahlbeck, Douglas PPLICANT: Katagiri, Fumiaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT: Staskawicz, Brian J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Lech, Karen F.
REGISTRATION NUMBER: 35,238
REFERENCE/DOCKET NUMBER: 00786/254001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEPAX: (617) 542-8906
TELEPAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
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FILING DATE: April 13, 1994
ATTORNEY_AGENT INFORMATION:
NAME: Lech, Karen F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30B CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/310,912A FILING DATE: September 22, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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ZIP: 02110-2904
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TOPOLOGY: linear
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  2.3%;
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                                                                  US-09-330-330-9
                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear; MOLECULE TYPE: protein PCT-US95-04589-190
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PCT-US95-04589-190
Sequence 9, Application US/09330330 Patent No. 6274789
GENERAL INFORMATION:
APPLICANT: Yano, Masahiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 190, Application PC/TUS9504589 GENERAL INFORMATION:
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                                                                                                                                                                                            Best Local Similarity Matches 7; Conserv
                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/04589
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (617) 542-890
TELEX: 100254
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/227,360
FILING DATE: 13-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02110-2904
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street Suite 3100
                                                                                                                             256 LALITLG 262
|||||||
5 LALITLG 11
                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 00:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEPAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yu, Guo-Liang
TITLE OF INVENTION: RPS2 GENE AND USES THEREOF
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Katagiri, Fumiaki
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Maximum DB seq length: 200000000
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Match
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Copyright (c) 1993 - 2000 Compugen
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AAB27272
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868, an A33 an
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AAB80409
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AAB838333
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AAB3838383
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AAY10191
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                                                                    ALIGNMENTS
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Arabidopsis thalia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunoglobulin superfamily; Ig Sf; vascular adhesion molinflammation; cancer; wound; angiogenesis; human; confluency regulated adhesion molecule 1; CRAM-1; JAM-2
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                                                                                                                                                                                                                                                                                                                                                                                             Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1) Polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability
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Pred. No. 1.3e-288;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin superfamily; Ig Sf; vascular adhes inflammation; cancer; wound; angiogenesis; mouse; confluency regulated adhesion molecule 1; CRAM-1;
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                                                                                                                                                                                                                        MALSRRLRLRYARLPHFFLLLLFRGCMIEAVNLKSSNRNPVVHEFESVELSCIITHSQT
NIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTSEEG
                                                                                                                                             NDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRNDVPL
                                                                                                                                                                             2000-587436/55
DB; AAA97189.
                      DFRHKSSFVI 310
                                                                                                 PTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEVYDL
                                                                                                                                   ndrkevdeitielivqvkpvtpvcripaavpvgktatlqcqesegyprphyswyrndvpl
                                                                                                                                                                                                    SDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEVVAL
                                          niagiiggvlvvlivlavitmgiccayrrgcfisskqdgesykspgkhdgvnyirtseeg
                                                                                     \verb|ptdsranprfqnssfhvnsetgtlvfnavhkddsgqyyclasndagaarcegqdmevydl|
                                                                                                                                                                                                                                                                       310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                d human Confluency Regulated Adhesion Molecule polypeptide, useful for treatment of tumors, in and modulating vascular permeability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RMF
                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                           310
                                                                                                                                                                                                                                                                                                                                                  of wound
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                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            8; 59pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99EP-0200746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein;
                                                                                                                                                                                                                                                                                                                                                healing
                                                                                                                                                                                                                                                                                100.0%;
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Pred. No. 1.3
); Mismatches
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                                                                                                                                                                                                                                                                                DB 21;
L.3e-288;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            adhesion molecule;
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                                                                                                                                                                                                                                                                                         Length
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L protein
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                                                                                                                                                                                                                                                                                                                                                                                                      molecule
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RESULT
AAB392L7
AAB392A
ID AAB3
XX AAB3
AC AAB3
XX AAB3
XX Huma
XX Huma
KW Huma
KW ant!
KW noot
KW opt!
KW Opt!
KW Opt!
KW Wour
XX Home
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC The polynucleotide sequences given in AAC74223-C74279 encode the human CC secreted proteins represented in AAB39179-B39226. Sequences AAB39227-B39308 are alternative proteins encoded by the genes, and also CC AAB39227-B39308 are alternative proteins encoded by the genes, and also CC protein sequences with which they share homology. The proteins have activities based on the tissues and cells in which they are expressed. CC Examples of activities include: immunosuppressive; antiarthritic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; anotropic; neuroprotective; antibacterial; virucide; fundicide; and opthalmalogical. The human secreted proteins, fundicide; and opthalmalogical. The human secreted proteins, complexed and opthalmalogical. The human secreted proteins of the invention may be useful complexed and conditions such as autoimmune diseases e.g. rheumatoid complexed and conditions such as autoimmune diseases e.g. rheumatoid complexed and conditions such as autoimmune diseases e.g. rheumatoid complexed and conditions such as autoimmune diseases e.g. rheumatoid complexed and conditions such as autoimmune diseases e.g. rheumatoid complexed expensive discorders e.g. cardiac arrest, cerebrovascular complexed expensive e
Query Match
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                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 32; 434pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-MAR-1999;
10-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-MAR-2000; 2000WO-US06792
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted protein sequence encoded by gene 15 SEQ ID NO:134.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cerebrovascular disorder; nervous system disorder; ocular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hyperproliterative disorder; cardiovascular disorder; infection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0125362
99US-0169980
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Komatsoulis G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antirheumatic;
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ABG04645
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                                                  to restore normal activity of (II) or to treat disease states involving (III). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifications.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 20; SEQ ID No 35004; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-639362/73.
N-PSDB; AAS68832.
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23-AUG-2000; 2000US-0649167
                                    at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drmanac RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAR-2001; 2001WO-US08631.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; chromosome mapping; gene mapping; gene therapy; forensic
food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human diagnostic protein #4636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG04645;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABG04645 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                for identifying expressed genes. (I) is useful in gene therapy
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  291 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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Matches
 167
PRPHYSWYRNDVPLPTDSRANPRF 190
                           Similarity
                 7.7%; ilarity 100.0%; Conservative
                 0;
                          Score 24;
Pred. No.
                  Mismatches
                          DB 21;
1.1e-14
                          .1e-14;
                                 Length 285;
                 Indels
                 0,
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Matches Query Match Best Local (

Local Similarity

100.0%;

Score 24; Pred. No. Mismatches

1.1e-14; DB 22;

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RESULT
AAB27276
ID AAB27276
AC AAB2
XX AAB2
XX Inmn
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KW Conil
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AAB33457
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Matches 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunoglobulin superfamily; Ig Sf; vascular adhesion molinflammation; cancer; wound; angiogenesis; human; confluency regulated adhesion molecule 1; CRAM-1; JAM-2.
  antianaemic;
         Human; immune related disease; diagnosis; antiinflammatory; cardian dermatological; antiarthritic; antirheumatic; immunosuppressive; haemostatic; antithyroid; antidiabetic; nootropic; neuroprotective;
                                                                                                                                                                                                                                                                      The present sequence is the human confluency regulated adhesion molecule I (CRAM-1, also known as JAM-2). CRAM-1 is one of the vascular adhesion proteins of the immunoglobulin superfamily (Ig Sf). The CRAM-1 protein and coding sequence can be used in the treatment of cancer, inflammation to modulate cell-cell interactions and angiogenesis, and in the
                                                                                                                                                                                                                                                                                                                                                 Isolated human Confluency Regulated Adhesion Molecule 1 or 2 (CRAM-1 CRAM-2) polypeptide, useful for treatment of tumors, inflammation reactions and modulating vascular permeability -
                                                  Human
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                                                                     29-JAN-2001
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                                                                                                                                                                                                                                                                                                                                 Claim
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                                                                                                           AAB33457
                                                                                                                                                                                                                                            Sequence
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DB; AAA95306.
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                                                PRO1868 protein
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24; Conservative
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                                                                                                          standard; Protein;
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                                                                                                                                                                                                                                                                                                                              6; 59pp; English.
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 hepatotropic;
                                                                   (first entry)
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  virucide;
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                                                                                                                                                                                             Score 24; DB; Pred. NO. 1.2
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                                                SEQ
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                                                 ID NO:193.
 antipsoriatic;
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1.2e-14
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es 0;
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 antiallergic;
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 05-JAN-2000;
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06-JAN-2000;
11-FEB-2000;
18-FEB-2000;
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22-FEB-2000;
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08-SEP-1999

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15-SEP-1999

05-OCT-1999

29-OCT-1999

29-NOV-1999

30-NOV-1999

30-NOV-1999

01-DEC-1999

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02-DEC-1999

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23-MAR-1999;
12-APR-1999;
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23-JUN-1999;
20-JUL-1999;
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10-MAR-1999;
12-MAR-1999;
 Claim 33; Fig
                                                                                                Ashkenazi
Kabakoff R
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14-MAY-1999;
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28-JUL-1999;
                                                                                      Stewart TA,
                                                                                                                             (GETH )
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DB; AAC58622.
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2000WO-US00277.
2000WO-US00376.
2000WO-US03565.
2000WO-US04341.
2000WO-US04342.
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88; 309pp; English
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99WO-US12252
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99WO-US20111
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99US-0162506.
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99WO-US28313.
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Lu Y,
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Van J, Pennica |
Watanabe CK, 1
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antiasthmatic; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; spondyloarthropathy; systemic sclerosis; sarcoidosis; idiopathic inflammatory myopathy; Sjogren's syndrome; thyroiditis; systemic vasculitis; autoimmune haemolytic anaemia; dabetes mellitus; autoimmune thrombocytopaenia; immune-mediated renal disease; demyelinating disease; hepatobiliary disease; Whipple's disease;
                                                                                                             inflammatory bowel disease; gluten-sensitive enteropathy; autoimmune disease; immune-mediated skin disease; allergic disease;
                                                                  immunological disease; transplantation associated disease;
graft-versus-host-disease.
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Gurney AL, Hebe D, Shelton DL, Wood WI, Yan M; D, Wood Hebert ert C, Smith Henzel

Σ

Sixty four PRO polypeptides, useful in the diagnosis and treatment immune related disorders, e.g. systemic lupus erythematosis, rheumarthritis, osteoarthritis, thyroiditis and diabetes mellitus rheumatoid . of

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RESULT
AAY96735
ID AAY9
XX
AC AAY9
AC AAY9
AC AAY9
CF PRO1
XX
DF PRO1
XX
PRO1
XX
PRO1
XX
OS HOMC
XX
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Best Local Similarity
Matches 24; Conserv
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                                                                   Modified-site
                                                                                                                                                                                                            Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRO1868; A33 antigen; secreted protein; transmembrane protein; anti-inflammatory; cytostatic; recombinant production; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRO1868, an A33 antigen homologue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY96735;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY96735 standard; Protein; 310 AA
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                                                                                                                                                                                                                                                                                   Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anaemia, autoimmune thrombocytopaenia, thyroiditis, diabetes mellitus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            167
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226..231
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215..220
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107..1
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69..77
                                                                   243..248
                                                                                                                                                                                                                                                                                                                                                   192..195
                                                                                                /label= Transmembrane_domain
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                      "N-myristoylation site"
                                                                                                                                                                   "N-myristoylation site"
                                                                                                                                                                                                                                      "N-myristoylation site"
                                                                                                                                                                                                                                                                                                             "N-glycosylation site"
                                                                                                                                                                                                                                                                                                                                                                                "cAMP- and cGMP-dependent phosphorylation site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Casein kinase II phosphorylation site"
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Pred. No.
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. 1.2e-14;
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RESULT
AAY96294
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12-JAN-1999;
12-JAN-1999;
12-JAN-1999;
09-FEB-1999;
10-FEB-1999;
12-FEB-1999;
                                                                                                                                                                                                                                                                                                                 This is PRO1868, a putative homologue of A33 antigen, a known colorectal cancer-associated marker. The invention concerns novel secreted and transmembrane proteins, designated PRO polypeptides. The cDNA and gene sequences are useful in the recombinant production of PRO polypeptides, as a hybridization probe to screen libraries to isolate cDNAs with sequence identity to PRO polypeptides or to map the gene encoding the PRO polypeptides and analyzing genetic disorders. The cDNA/gene can also be used to produce transgenic animals useful for the development and screening of therapeutically useful reagents. They can also be used in gene therapy, e.g. to replace a defective gene.
          Human; immunoglobulin; IGFAM-6; IGFAM; immune disorder; cancer; infection; inflammation; haematopoiesis; AIDS; allergy.
                                              Human IGFAM-6 immunoglobulin.
                                                                        16-AUG-2000
                                                                                               AAY96294;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Fig 14; 154pp; English.
                                                                                                                     AAY96294 standard; protein; 310
                                                                                                                                                                                                                                                                                              Sequence
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N-PSDB; AAA51265.
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16-DEC-1998;
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                                                                                                                                                                              transmembrane polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated nucleic acid molecule encodes a
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                                                                                                                                                                                                                                                                                              310 AA;
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                                                                      (first entry)
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Pan J, Roy MA,
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990S-011558.
990S-0115565.
990S-0115733.
990S-0119341.
990S-0119357.
990S-0119357.
990S-0119352.
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98US-0113145
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296..299
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, Stewart TA,
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                                                                                                                                                                                                                                                        DB 21;
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                                                                                                                                                                                                                                                       Length 310;
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D, Watanabe
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Homo sapiens

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RESULT
AAM93323
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                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                      Matches
                                                                                                                                                                                                                                           tissue. It is expressed in reproductive, nervous and cardiovascular tissue, where cancer and inflammation are common. The gene, protein, its antibodies, agonists and antagonists are suitable for diagnosing and treating many diseases, including cancer, immune system disorders (such as inflammation, AIDS, allergies, anaemia, arteriosclerosis, asthma, atherosclerosis, cholecystitis, Crohn's disease, diabetes mellitus, emphysema, Graves' disease, hepatitis, multiple sclerosis, psoriasis, rheumatoid arthritis, scleroderma, systemic lupus erythematosus and ulcerative colitis), complications of cancer, haemodialysis and extracorporeal circulation, trauma and haematopoietic cancer (such as leukaemia) and infections caused by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-NOV-1998;
22-DEC-1998;
07-APR-1999;
06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                           The
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin superfamily proteins, the agon: protein is useful for preventing and treating altered levels of the protein such as cancer,
                       AAM93323;
                                             AAM93323 standard;
                                                                                                                                                                                                                                                                                                                                                                  IGFAM-6. Its gene was is tissue. It is expressed
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                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 82-83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB;
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                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                      bacteria,
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                                                                                                                 167 PRPHYSWYRNDVPLPTDSRANPRF 190
                                                                                                       167
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                                                                                                                                                                                                                                                                                                                                                                                          present sequence is the human immunoglobulin superfamily
                                                                                                   prphyswyrndvplptdsranprf 190
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Lal
                                                                                                                                                   Similarity 100.
24; Conservative
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                                                                                                                                                                                                                                                                                                                                                                               Its gene was isolated from a cDNA library
                                                                                                                                                                                                                                     viruses,
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(first
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98US-0113635.
99US-0128194.
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238..260
/label= t
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153..221
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46..117
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31..310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Corley NC,
Hillman JL,
                                             Protein;
                                                                                                                                                                                                                                     fungi or parasites.
                                                                                                                                                                7.7%;
entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 105pp; English.
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                                             310
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                                                                                                                                                                Score 24;
Pred. No.
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Yang J;
                                              A
                                                                                                                                                     Mismatches
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treating disorders associated wis
cancer, immune system disorders
                                                                                                                                                                 1.2e-14;
                                                                                                                                                                           DB 21;
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AAM93905
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Best Local Similarity 100.
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                          clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a polypeptide encoded by a full length human cDNA encoded by a full length human cDNA encoded by a full length human cDNA for the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in CD-ROM format directly from EPO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     830 Primers useful for synthesizing use in genetic manipulation -
                                EP1130094-A2
                                                                            Human;
                                                                                                   Human
                                                                                                                          06-NOV-2001
                                                                                                                                                 AAM93905;
                                                                                                                                                                         AAM93905 standard;
                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to primers for synthesising full length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wakamatsu A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUL-2000; 2000EP-0114089
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                                                     lomo sapiens
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                                                                                                 polypeptide,
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                                                                            full
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK94243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO
                                                                                                                                                                                                                                                                                                                                        310 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; 99JP-0194486.
; 2000JP-0118774.
; 2000JP-0183765.
                                                                            length
                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sugiyama
                                                                            CDNA;
                                                                                                   SEQ ID NO:
                                                                                                                                                                         Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2845; 1380pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA;
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                                                                                                                                                                                                                                                                                           100.0%;
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Nagai
                                                                           cDNA synthesis;
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                                                                                                                                                                         310
                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                         Score 24;
Pred. No.
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K, Kojima S,
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1.2e-14;
                                                                            oligo-capping
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Otsuki T, Koga
                                                                                                                                                                                                                                                                                                     Length
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05-SEP-2001

07-JUL-2000; 2000EP-0114089

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RESULT 1
AAU12440
ID AAU1
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Best Local S
Matches 24
01-DEC-1999;
01-DEC-1999;
02-DEC-1999;
02-DEC-1999;
02-DEC-1999;
02-DEC-1999;
09-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a polypeptide encoded by a full length human cDNA of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in CD-ROM format directly from EPO.
                                                                                                                                                                                                           Human secretory and transmembrane; PRO; mammalian; cancer; lung; breast; prostate; cervical; tumour necrosis factor-alpha; TNF-alpha; cartilage; ear; proliferation; glucose; free fatty acid; skeletal muscle; adipocyte; A-peptide; factor VIIA; gene therapy.
                                                                                                 01-DEC-2000;
                                                                                                                              07-JUN-2001.
                                                                                                                                                       WO200140466-A2
                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                    Human PRO1868
                                                                                                                                                                                                                                                                                                               24-OCT-2001
                                                                                                                                                                                                                                                                                                                                           AAU12440
                                                                                                                                                                                                                                                                                                                                                                     AAU12440 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JUL-1999; 99JP-0194486
11-JAN-2000; 2000JP-0118774
02-MAY-2000; 2000JP-0183765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      830 Primers useful for synthesizing use in genetic manipulation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wakamatsu
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Similarity 100.0%;
24; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  310
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                  2000WO-US32678
                                                                                                                                                                                                                                                                                  polypeptide sequence
              99WO-US28301.
99WO-US28634.
99WO-US28551.
99WO-US28564.
99WO-US28565.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ĄĄ;
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a T, Nagai K, Kojima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24;
; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 22; I
1.2e-14;
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S, Otsuki T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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167

Query Match Best Local S Matches 24

Similarity 100 24; Conservative

100.0%;

; Score 24; DB %; Pred. No. 1.2 0; Mismatches

1.2e-14; hes 0; DB 22;

Indels

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Gaps

0

Length 310;

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167

RESULT 12 AAB80272

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CC breast, prostate, rectal, cervical or liver tumours by comparing PRO CC polypeptide expression in a cell sample to that in a control sample. CC Some of the 275 sequences are also useful to stimulate the release of CC tumour necrosis factor-alpha (TNF-alpha) from human blood, the CC proliferation or differentiation of chondrocytes, the proliferation or cells, the release of proteoglycans from CC cartilage, the proliferation of inner ear utricular supporting cells or CC cartilage, the release of a cytokine from peripheral blood CC monocytes (PBMCs), or the proliferation of endothelial cells. Some of CC concepts the cells or by adipocytes; or inhibit binding of A-peptide cells or by adipocytes; or inhibit binding of A-peptide conceptues involved in binding interactions. The polypucleotides encoding CC molecules involved in binding interactions. The polypucleotides encoding CC pRO polypeptides can be used to generate probes, antisense RNA/DNA, CC transgenic or knock out animals and can be used in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Baker KP,
Gerritsen
Smith V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-FEB-2000;
24-FEB-2000;
01-MAR-2000;
20-MAR-2000;
21-MAR-2000;
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11-FEB-2000;
18-FEB-2000;
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20-DEC-1999;
20-DEC-1999;
30-DEC-1999;
06-JAN-2000;
Sequence
                                                                                                                                                                                                                                                                                                      AAU12172-AAU12446 represent novel human secretory and transmembrane PRO polypeptides. The PRO polypeptides are useful to detect other PRO polypeptides, to link bloactive molecules to cells expressing PRO polypeptides, to modulate biological activities of cells expressing PRO polypeptides, and to detect the presence of mammalian lung, colon, PRO polypeptides, and to detect the presence of mammalian lung, colon, PRO polypeptides, and to detect the presence of mammalian lung, colon, PRO polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Fig 538; 813pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated, secretory and transmembrane PRO polypeptide used to detect other PRO polypeptides, link bioactive molecules to cells expressing PRO polypeptides, and detect the presence of mammalian tumours e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; AAS21512.
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17-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      breast,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stewart TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beresini M, Deforge L, Desnoyers L, Filva:
ME, Goddard A, Godowski PJ, Gurney AL, Shv
Stewart TA, Tumas D, Watanabe CK, Wood WI,
 310 AA;
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99WO-US30991
99WO-US31243
2000WO-US0376
2000WO-US0376
2000WO-US03365
2000WO-US03414
2000WO-US04414
2000WO-US04414
2000WO-US05601
                                                                                                                                                                                                                                                                                                                                                                                                                                                      prostate, cervical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Filvaroff E, Gao
AL, Sherwood S;
Zhang Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gao
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Best Local Similarity 100 Matches 24; Conservative

100.0%;

Pred. No.

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Mismatches

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Indels

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Filvarum
Godowski PJ, Gra
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08-SEP-1999;
13-SEP-1999;
15-SEP-1999;
15-SEP-1999;
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29-NOV-1999;
30-NOV-1999;
16-DEC-1999;
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                                   Parkinson's disease), wound repair, cardiovascular disorders (e.g. endometrial bleeding angiogenesis, ischaemias such as coronary ischaemia, atherosclerosis), inflammatory disorders (e.g. asthma, rheumatoid arthritis, multiple sclerosis), infertility, AIDS and diabetes and retinal disorders such as retinitis pigmentosum. The PRO nucleic acids have applications in molecular biology, including use as hybridization probes, and in chromosome and gene mapping.
                                                                                                                                                                                                     transmembrane PRO polypeptides. The PRO polypeptides are useful for treating skin diseases (e.g. psoridasis), cancers (e.g. lung squamous cell carcinoma), gastrointestinal disorders (e.g. enterocolitis), neurodegenerative diseases (e.g. Alzheimer's disease, enterocolitis),
Sequence
                                                                                                                                                                                                                                                                                                  The present sequence is one of sixty
                                                                                                                                                                                                                                                                                                                                             Claim 1; Fig 124; 393pp;
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26-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                               squamous cell
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Filvaroff E,
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                                                                                                                                                                                                                                                                                                                                                                                            Alzheimer's
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                                                                                                                                                                                                                                                                                                                                                                                                             ty one nucleic acids encoding PRO polypeptides which are useful treatment of skin diseases (e.g. psoriasis), cancers (e.g. lung amous cell carcinoma) and neurodegenerative diseases (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-081051/09
DB; AAF72433.
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  310
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Fong S, Gao W, Gerber H,
Grimaldi CJ, Gurney AL, I
an J, Paoni NF, Roy MA, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US30095.
99WO-US30911.
99WO-US30999.
99WO-US00219.
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99WO-US28214.
99WO-US28313.
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99WO-US20594.
99WO-US20944.
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99WO-US21547.
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99US-0145698.
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                                                                                                                                                                                                                                                                                                                                             English.
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                                                                                                                                                                                                                                                                                                  one novel secreted
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Hillan KJ,
Stewart TA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ME, Goddard
Kljavin IJ;
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Query Match

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Length

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24-APR-2001

(first entry)

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RESULT 1
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                                                                                                                                                                                                      arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases
                                                                                                                                                                                                                                                                                                            Claim 11; Page 538-539; 601pp; English.
                                                                                                                                                                                                                                                                                                                                   and microorganism infections
                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200107459-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Secreted protein; human; autoimmune; hyperproliferation; cardiovascular; cerebrovascular; infection; food.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Secreted
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AAB80408
                     AAB80408
                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                          The present invention relates to 29 human secreted invention is used to prevent autoimmune diseases e.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-2000; 2000WO-US19735
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                                                                               167
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                                                                                                   167 PRPHYSWYRNDVPLPTDSRANPRF 190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRPHYSWYRNDVPLPTDSRANPRF 190
                                                                          prphyswyrndvplptdsranprf 190
                                                                                                                                                                                                                                                                                                                                                                                          2001-123261/13
                                                                                                                                                                                                                                                                                                                                                                                                                CA, Ruben SM,
, Lafleur DW,
                                                                                                                          Similarity
24; Conserv
                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein
                                                                                                                                                                                  310 AA;
                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                    protein;
                                                                                                                                   7.7%;
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Olsen HS,
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Pred. No.
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Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B
                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #13
                                                                                                                                   DB 22; 1
1.2e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                Ni J, Soppet DR, Komatsoulis GA;
                                                                                                                                              Length 310;
                                                                                                                          Indels
                                                                                                                                                                                                                                                                           proteins. The
                                                                                                                                                                                                                                                                                                                                               or disorders
                                                                                                                                                                                                                                                                                                                                                                     for
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                                                                                                                                                                                                                                                                                                                                                                                                                            Moore PA;
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                                                                                                                          0;
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Secreted

protein

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AAAB80409
ID AAB8
XX AAB8
AC AAB8
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AC Sec:
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Best Local S
Matches 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to 29 human secreted proteins. Th invention is used to prevent autoimmune diseases e.g. rheumatoi arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. cerebral ischemia, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid encoding 29 secreted proteins, diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                              20-JUL-2000;
                                                                                             01-FEB-2001
                                                                                                                                         WO200107459-A1
                                                                                                                                                                                          HOMO
                                                                                                                                                                                                                                      cardiovascular;
                                                                                                                                                                                                                                                          Secreted protein;
                                                                                                                                                                                                                                                                                                         Secreted protein
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                                                                                                                                                                                                                                                                                                                                                                                                         AAB80409
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB80409 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 11; Page 557-558; 601pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Secreted protein; cardiovascular; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    167 PRPHYSWYRNDVPLPTDSRANPRF 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                microorganism infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            prphyswyrndvplptdsranprf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-123261/13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 100.0%;
24; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lafleur DW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruben
                                              2000WO-US19735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US19735
                                                                                                                                                                                                                                                                                                                                                       (first entry)
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  99US-0145220
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                                                                                                                                                                                                                                      cerebrovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , MS
                                                                                                                                                                                                                                                                                                           encoded by
                                                                                                                                                                                                                                                        human; autoimmune; hyperproliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ebner R,
Olsen HS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       γď
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                                                                                                                                                                                                                                                                                                         gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                       310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Duan RD,
Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                           #39
                                                                                                                                                                                                                                      infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ine; hyperproliferation;
infection; food.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.2e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ni J, Soppet DR,
Komatsoulis GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proteins. The g. rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Moore PA;
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RESULT 1
AAB38333
ID AAB38333
ID AAB38333
AC AAB3
AC AAB3
AC AAB3
AC Immu
KW Cytc
KW Immu
KW Cytc
KW NOO1
KW VULI
XX
VOS HOME
COS HOME
COS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nootropic; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to 29 human secreted proteins. Th invention is used to prevent autoimmune diseases e.g. rheumatoi arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebrovas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders and microorganism infections
                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1999;
26-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; neuroprotectiv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g
                                                                                                                              WPI; 2000-647418/62.
                                                                                                                                                                                                                                                      Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200061623-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB38333;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      corneal infection. Also used in food preparations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; Page 559-560; 601pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-123261/13.
                                                                                                                                                                                                                        Lafleur DW,
                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRPHYSWYRNDVPLPTDSRANPRF 190
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, Lafleur I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibacterial; virucide; gene therapy; infection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          310 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                            Olsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000WO-US08979
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                                                                                                                                                                                                                                                                                                                      GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                     99US-0128693.
99US-0130991.
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                                                                                                                                                                                                                                                      Komatsoulis
                                                                                                                                                                                                                            HS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.7%;
100.0%;
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Olsen F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SCI INC.
                                                                                                                                                                                                                            Ebner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                r R,
HS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 24;
Pred. No.
                                                                                                                                                                                                                        R, GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ьy
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Birse CE,
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New nucleic acid molecules encoding 62 human secreted proteins diagnosing, preventing, treating or ameliorating medical condit

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RESULT 17
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            Claim 11; Page 642-643;
                                         diagnosing, preventing, used as food additives
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26-APR-1999;
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Florence KA,
                                                      62 human secreted proteins for ameliorating medical conditions
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Best Local :
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Lafleur DW,
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                                                                                                                                                                                              New nucleic acid molecules encoding diagnosing, preventing, treating or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 311
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Sequences AAB38321-B38396 represent the amino acid sequences of 62 human secreted proteins encoded by the genes AAC69512-C69587. The and proteins are useful for preventing, ameliorating or treating me

medical genes Claim

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Page 643-644; 716pp; English.

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                                                                                   The present invention relates to 29 human secreted proteins. The invention is used to prevent autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene
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cerebrovascular disorders e.g. cerebral ischemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. Also used in food preparations.
                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                      New isolated nucleic acid encoding 29 secreted proteins, for diagnosing, preventing and treating e.g. autoimmune, hyperproliferative, cardiovascular, and ocular diseases or disorders
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cerebrovascular; infection; food.
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Komatsoulis
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                                           Note: The sequence data for this patent did not specification, but was obtained in electronic for
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23-AUG-2000;
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food supplement; medical imaging; diagnostic;
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stic; genetic disorder.
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                                                                                                                     polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity and to produce other traits to assess biodiversity
                   and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention.

Note: The sequence data for this patent did not appear in the printer specification, but was obtained in electronic format directly from W
at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 20;
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medical imaging; diagnostic; genetic disorder.
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Matches 21; Conservative
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                                                 Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WI
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medical imaging; diagnostic; genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 21;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 22;
1e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 361;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mutations
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                                                   WIPO
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DB 22;

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RESULT 2
AAY11472
ID AAY1
XX
AC AAY1
DT 21-:
XX
DE Huma
XX
Huma
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Best Local S
Matches 15
Ouery Match

4.8%; Score 15; DB

Best Local Similarity 100.0%; Pred. No. 1.

Matches 15; Conservative 0; Mismatches
                                                                                                              sequences can be used for producing secreted human gene products. They can also be used to develop products for diagnosis and therapy. The proteins obtained may have cytokine activity, cell proliferation/differentiation activity, haematopolesis regulating activity, tissue growth regulating activity, reproductive hormone regulating activity, chemotactic/ chemokinetic activity, haematory activity activity, receptor/ ligand activity, anti-inflammatory activity tumour inhibition activity or other activities. The products can be used in forensic, gene therapy and chromosome mapping procedures. The sequences are also be used for obtaining corresponding promoter sequences. The nucleic acids encoding the signal peptide can be used for directing extracellular secretion of a polypeptide or the insertion of a polypeptide into a cell.
                                                                                                                                                                                                                                                                                                                                     human secreted proteins, and encode the proteins given in AAY11374 AAY11531, respectively. The proteins given represent the signal per and an N-terminal fragment of a secreted protein. The nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acids encoding human secreted - proteins obtained from cDNA libraries prepared from substantia nigra, cerebellum, surrena and fetal brain tissue
                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                     AAX39440 to AAX39597 represent 5' expressed sequence tags (ESTs) for human secreted proteins, and encode the proteins given in AAY11374 to
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 34; Page 394; 434pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-153781/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Duclert A, Dumas Milne Edwards J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W09906551-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                forensic; gene therapy; chromosome mapping; signal peptide; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopolesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY11472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY11472 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; secreted protein; EST; expressed sequence tag; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    thrombolytic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sqtsdpriewkkiqd 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SQTSDPRIEWKKIQD 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST
                                                                                      89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                      ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-inflammatory; tumour inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0905133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-IB01235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.88;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Š
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                 DB 20; I
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                                 Length 89;
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RESULT 2
ABG27038
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                                                                                                          cc polypeptide (II) sequences. (I) is useful as hybridisation probes, cc polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The cand gene mapping, and in recombinant production of (II). The complex com
                                           Note: The sequence data for this patent did not appear in the specification, but was obtained in electronic format directly at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 20; SEQ ID No 57397; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to isolated polynucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         biodiversity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-639362/73
N-PSDB; AAS91225.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Drmanac RT, Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAR-2001; 2001WO-US08631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; chromosome mapping; gene mapping; gene therapy; forensic
food supplement; medical imaging; diagnostic; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG27038;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sqtsdpriewkkiqd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
  267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0540217.
2000US-0649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72
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                                                                    from WIPO
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Query Match Best Local Similarity

Matches

10;

Conservative

0

100.0%;

Score 10; Pred. No. Mismatches

DB 22; 0.27; 0

Length 267 Indels

0;

Gaps

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RESULT :
AAWOILD ID
AAWOILD AAW
XX
XX
AC AAW
XX
DT 15-N
DX 5-1
DX Flee
KW Flee
KW Conni
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                    release formulations to reduce the flea burden on and around the animal. The inhibitors may be included in insecticidal compositions to increase efficacy of other active agents, by reducing proteolytic activity in the flea midgut.
                                                                              library by homology with conserved serine protease sequences. The sequence contains part of N-terminal sequence AAW01177, and other SP12 sequences are given in AAW01197 (C-terminal sequence) and AAW01209 (PfSP12-69). The peptide may be used in a vaccine for protection of domestic animals from flea infestation, or in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Flea;
                                                                                                                                          This sequence represents a partial N-terminal sequence of midgut serine protease, Spl2-52, isolated from a flea cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Siphonaptera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               controlled release formulation; synergist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Serine protease PfSP12-52 N-terminal fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW01191;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW01191 standard; Peptide;
Sequence
                                                                                                                                                                            Claim 71; Page 163;
                                                                                                                                                                                                  DNA encoding Flea serine protease and vaccines to protect animals from flea
                                                                                                                                                                                                                                                                                                                                                                                                                                       25-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                              W09611706-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-MAR-1997
                                                                                                                                                                                                                                     WPI; 1996-221762/22
N-PSDB; AAT40836.
                                                                                                                                                                                                                                                                                  Arfsten
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                                                                                                                                                                                                                                                                                                                                                       07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                18-OCT-1995;
                                                                                                                                                                                                                                                                       Hunter
                                                                                                                                                                                                                                                                                                        (HESK-)
                                                                                                                                                                                                                                                                                                                     (PARA-)
                                                                                                                                                                                                                                                                                                                                                                             18-OCT-1994;
                                                                    solation of protease-inhibitors, which may be used in controlled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      midgut;
                                                                                                                                                                                                                                                                       n A,
                                                                                                                                                                                                                                                                                                        PARAVAX INC.
HESKA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           agut; serine protease;
animal; infestation;
48
                                                                                                                                                                                                                                                                      Dale B,
Rushlow
                                                                                                                                                                                                                                                                                 Dale
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AA;
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94US-0326773.
95US-0482130.
95US-0484211.
95US-0485443.
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/note=
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20..48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
1..19
                                                                                                                                                                           241pp; English.
                                                                                                                                                                                                                                                                       KΕ,
                                                                                                                                                                                                                                                                      Frank GR, Grie
KE, Stiegler GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Part of N-terminal fragment AAW01177"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          insecticide; protease-inhibitor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PfSP12-52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 À
                                                                                                                                                                                                                                                                                 Grieve RB,
                                                                                                                                                                                                  aminopeptidase
infestation.
                                                                                                                                                                                                                                                                       Yamanaka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vaccine;
                                                                                                                                                                                                               •
                                                                                                                                                                                                               useful in
                                                                                                                                                      flea
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Query Match Best Local S Matches 8

Local Similarity nes 8; Conserv

Conservative

2.6%; Score 8; DB: 100.0%; Pred. No. 5; tive 0; Mismatches

17;

Length 48; Indels

0

0

Gaps

0

Query Match Best Local Similarity

2.6%;

Score Pred.

8; No.

DB 5; 22;

Length 48;

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В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
                                       protease activity comprising: (a) contacting (1) with a protease substrate and a putative inhibitory compound, where (I) has proteolytic activity in the absence of the compound; and (b) determining if the compound inhibits protease activity by detecting cleavage of the protease substrates; where decreased cleavage of the protease substrate indicates an inhibitory compound. (I), nucleic acid molecules encoding (I), and antibodies immunospecific for (I) and (II) are useful for preventing and reducing flea infestations, particularly the species Ctenocephalides felis (cat fleas). Ctenocephalides canis (dog fleas) and pulex irritans, in animals, preferably cats and dogs. They are also useful for reducing infestation by other ectoparasites, preferably mosquitoes, midges, sandflies, blackflies, ticks and Rhodnius. AAC90818 to AAC90913 and AAB50551 to AAB50644 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1995;
07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-DEC-1991;
18-OCT-1994;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                       New isolated flea proteins with proteolytic activity, useful preventing and reducing flea infestations in mammals especial and dogs \dot{\cdot}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Flea; cat flea; serine protease; aminopeptidase; cysteine pro
flea infestation; proteolytic; insecticide; vaccine; cat; dog
Ctenocephalides felis; Ctenocephalides canis; Pulex irritans;
                                                                                                                                                                                                                                  The present invention describes isolated flea serine protease, aminopeptidase and cysteine protease proteins (I). Also described method for identifying a compound (II) capable of inhibiting flea
   Sequence
                              exemplification of
                                                                                                                                                                                                                                                                                             Claim 1; Column 47-48; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-136374/14.
N-PSDB; AAC90844.
                                                                                                                                                                                                                                                                                                                                                                                                                             Grieve
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US6150125-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ctenocephalides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ectoparasite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Flea serine protease nfSP12 N-terminal protein sequence #65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB50588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB50588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HESK-)
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                                                                                                                                                                                                                                                                                                                                                                                                                             PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HESKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; Protein;
   48
                                                                                                                                                                                                                                                                                                                                                                                                                                          Silver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORP.
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95US-0484211.
95US-0485443.
95US-0485455.
97WO-US14442.
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94US-0326773
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                                                                                                                                                                                                                                                                                                                                                                                                                                          ç,
                            the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Rushlow KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Hunter SW,
                                                                                                                                                                                                                                                                                                                                                                                                                                          Frank GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cysteine protease;
lne; cat; dog; dog
                                                                                                                                                                                                                                                                                                                                      especially cats
                                                                                                                                                                                                                                                                                                                                                                                                                                          Stiegler GL;
                                                                                                                                                                                                                                                                                                                                                       for
                                                                                                                                                                                                                                                     s,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                flea;
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                                                                                                                                                                                          AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AAG81454 to AAG83120, from Staphylococcus epidermidis. (I) and (II) can have antibacterial activity and therefore can be used (I) and (II) can have antibacterial activity and therefore can be used (I) vaccination. The nucleic acids (I) may be used to produce the CC s. epidermidis polypeptides (II) via the production of vectors (I) epidermidis polypeptides (II) via the production of vectors (II) via the production of vectors (II) epidermidis polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their activity and therefore identify compounds that may be used for the CC treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA (I) epidermidis sequences from the present invention. AAH53971 to AAH55090 represent oligonucleotide sequences and primers which are used (I) the exemplification of the present invention. The polynucleotide sequences siven in the sequence listing of the present specification, the present specification, and the sequence listing only goes up to SEQ ID NO:4454 so even though sequences are given in the disclosure for SEQ ID NO:4465 to 4472,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
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                                                                   Matches
                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Staphylococcus epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Staphylococcus épidermidis SR1 strain; infection; diagnosis; vaccination; endocarditis.
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB; AAH52509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kimmerly WJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-NOV-2000; 2000WO-US30782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG81659;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAG81659 standard; Protein;
                                                                                                                                                                                 no sequences are present for SEQ ID NO:4455 to 4464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GLAX ) GLAXO
 181
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1 vlivlavi 8
                                242 IAGIIGGV 249
                                                                 Local Similarity
nes 8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27
iagiiggv 188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 151; 2188pp; English.
                                                                                                                                                    204
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                               2.6%;
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                                                                                  Score 8;
Pred. No.
                                                                 Mismatches
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                                                                        DB
18;
                                                                                                22;
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                                                                                              Length 204;
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                                                               Gaps
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AAW41974 ID AAW4

AAW41974 standard; Protein; 246

RESULT

29

Ър

181

iagiiggv 188

242 IAGIIGGV 249

Query Match Best Local s Matches 8

Similarity 100 8; Conservative

2.6%;

Score 8; DB 2; Pred. No. 18; 0; Mismatches

DB . 18; 22; 0

Length 204; Indels

0

Gaps

0

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RESULT
AAG82062
                             activity and therefore identify compounds that may be used for the treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA polynucleotide sequences from the present invention. AAH55091 to AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention.

N.B. The present invention specifically claims all the polynucleotide sequences given in the sequence listing of the present specification, however the sequence listing only goes up to SEO ID NO:4454 so even though sequences are given in the disclosure for SEO ID NO:4465 to 4472, no sequences are present for SEO ID NO:4465 to 4472.
                                                                                                                                                                                                                                        in vaccination. The nucleic acids (I) may be used to produce the S. epidermidis polypeptides (II) via the production of vectors containing them which are used to produce hosts cells which express the polypeptides. The polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their the polypeptides may also be used to assay for other inhibitors of their
                                                                                                                                                                                                                                                                                                                                                          AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AAG81454 to AAG83120, from Staphylococcus epidermidis (I) and (II) can have antibacterial activity and therefore can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 18; Page 349; 2188pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Staphylococcus epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S. epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-SEP-2001
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-316495/33.
N-PSDB; AAH52912.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kimmerly WJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vaccination;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Staphylococcus epidermidis SR1 strain; infection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAG82062;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAG82062 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GLAX ) GLAXO GROUP LTD
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204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  endocarditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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Ŗ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          204
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ΧIJ
                         RESULT 30
AAB50616
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                                                                                                                                                                                                        The present sequence represents a novel flea serine protease. The CC protease, its mimetopes, antibodies (Ab) and inhibitors of the CC protein, as well as the DNA encoding the protein, may all be used in the protein, as well as the DNA encoding the protease activity (especially in CC immunoglobulin protease) and so reduce flea protease activity (especially in CC cats or dogs. Alternatively, flea larvae may ingest the faeces of adult fleas which comprises anti-protease antibodies produced by a host animal in response to administration of the protein. Therapeutic compositions may further comprise a compound that reduces haematophagous ectoparasite burden by a method other than by reducing flea immunoglobulin protease activity. The novel flea DNA encoding the protein can also used to produce recombinant protein, and fragments of it are used as probes and CC primers for identification and isolation of related sequences, also as comparises to fite protein. Ab are also useful for screening expression of the protein and to target cyrotypes.
                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-APR-1997;
24-APR-1996;
15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Frank GR,
Silver G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Flea
            AAB50616 standard; Protein; 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Siphonaptera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Flea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New flea protease genes for the prophylaxis and cats or dogs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9740058-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 260-261; 318pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-OCT-1997
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                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HESK-) HESKA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunoglobulin
                                                                                                                                                                                                                                  libraries,
                                                                                                         252 VLIVLAVI 259
                                                                                                                                                 Local Similarity
                                                                               1998-076762/07.
DB; AAV04597.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              serine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Siphonaptera; protease; vaccine; prophylaxis; infestation; oglobulin protease; larvae; host animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaines PJ,
Stiegler GI
                                                                                                                                                                                                        246
                                                                                                                                                                                                                                 to purify the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protease SEQ ID
                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97US-0042945.
96US-0639075.
96US-0749699.
                                                                                                                                                                                                        AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GL;
                                                                                                                                 2.6%; Score 8; 1
100.0%; Pred. No.
Live 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Grieve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded by TAA, a stop codon"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and proteins treatment of
                                                                                                                                                                                                                                 protein and to target cytotoxins to fleas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NO:129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RВ,
            Ā
                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hunter SW,

    used in vaccine compositions
flea infestation, especially in

                                                                                                                                          DB .
21;
                                                                                                                                                              19;
                                                                                                                                   0;
                                                                                                                                                             Length 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rushlow
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                                                                                                                                   0;
                                                                                                                                   Gaps
                                                                                                                                   0;
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RESULT 31 AAG93112

В δÃ

N

vlivlavi

9

Query Match
Best Local Similarity
Matches 8; Conser

Conservative

0

2.6%;

Score 8; DB 2; Pred. No. 21; 0; Mismatches

DB 21; 22;

0, Length

Indels

0;

Gaps

0

252 VLIVLAVI 259

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CC aminopeptidase and cysteine protease proteins (I). Also described is a CC method for identifying a compound (II) capable of inhibiting flea CC protease activity comprising: (a) contacting (I) with a protease CC substrate and a putative inhibitory compound, where (I) has protease CC compound inhibits protease activity by detecting cleavage of the CC compound inhibits protease activity by detecting cleavage of the CC protease substrates; where decreased cleavage of the protease substrate (I), and antibodies immunospecific for (I) and (II) are useful for CC preventing and reducing flea infestations, particularly the species CC tenocephalides felis (cat fleas), Ctenocephalides canis (dog fleas) and CC pulex irritans, in animals, preferably cats and dogs. They are also CC useful for reducing infestation by other ectoparasites, preferably CC mosquitoes, midges, sandflies, blackflies, ticks and Rhodnius. AAC90818 CC targetic constitution of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-DEC-1991;
18-OCT-1994;
07-JUN-1995;
07-JUN-1995;
07-JUN-1995;
07-JUN-1995;
15-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Flea; cat flea; serine protease; aminopeptidase; cysteine protease; flea infestation; proteolytic; insecticide; vaccine; cat; dog; dog Ctenocephalides felis; Ctenocephalides canis; Pulex irritans;
                                                                                                                                                                                                                                                                                                       New isolated flea proteins with proteolytic activity, preventing and reducing flea infestations in mammals and dogs -
                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-136374/14
N-PSDB; AAC90878.
                                                                                                                                                                                                                                                                                                                                                                                                   Gaines
Grieve
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94US-0326773.
95US-0482130.
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Title:
Perfect score:
Sequence:
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length: 2000000000
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/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
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                US-08-485-455D-65
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US-08-906-769-65
US-08-906-769-65
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US-08-906-616-65
US-08-817-795-65
US-08-817-431-65
US-09-012-431-65
US-09-012-692-65
US-09-018-906-613-65
US-08-906-769-127
US-08-906-616-127
US-08-906-616-127
US-08-906-613-27
US-09-012-431-127
US-09-012-431-127
US-09-013-813-127
US-09-13-08-96-13-127
US-09-13-08-96-13-127
US-09-13-08-91-13-13-15
US-09-13-08-91-13-15
US-09-13-13-16
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US-08-861-464-8
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Sequence 65, ...
Sequence 65, ...
Patent No. 5712143
Patent No. 5712143
GENERAL INFORMATION:
APPLICANT: Grieve, Robert B.
APPLICANT: Rushlow, Keith E.
APPLICANT: Wu Hunter, Shirley
APPLICANT: Stiegler, Gary L.
TITLE OF INVENTION: FLEA PROTEINS, NUCLEIC ACID
TITLE OF INVENTION: MOLECULES, AND USES THEREOF
NUMBER OF SEQUENCES: 79
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                 RESULT 1
US-08-485-455D-65
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US-09-540-245A-15
US-08-685-589A-164
US-07-960-510-6
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US-07-960-510-7
S171673-7
US-07-960-510-7
S171673-7
US-09-053-197A-72
US-09-053-197A-72
US-09-053-197A-77
US-09-053-14-095D-18
US-08-672-345C-18
US-08-672-345C-18
US-08-672-345C-18
US-08-672-345C-18
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US-08-214-095D-98
US-08-214-095D-98
US-08-214-095D-98
US-08-22-573-2
US-08-441-629-13
US-08-441-629-13
US-08-441-629-13
US-08-441-495-27
US-08-441-495-27
US-08-441-495-27
US-08-461-91
US-08-06-616-91
US-08-06-769-91
US-08-06-769-91
US-08-06-613-91
US-08-06-613-91
US-08-07-044-1
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US-08-462-169B-13
US-08-462-169B-13
US-08-463-907-043-1
US-08-463-908-17-07-13
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TOPOLOGY: linear; MOLECULE TYPE: protein US-08-485-455D-65
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ATTORNEY/AGENT INFORMATION:
NAME: CONNELL, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE, DOCKET NUMBER: 2618
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
TELEPAX: (303) 863-0223
INFORMATION FOR SEQ ID NO: 65:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
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Best Local Similarity
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TELEFAX: (303) 863-0223
INFORMATION FOR SEO ID NO: 65:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                   SOFTWARE: PatentIn Release CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
CLASSIFICATION:
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REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Stiegler, Gary L.
TITLE OF INVENTION: FLEA PROTEASE PROTEINS, NUCLEIC ACID
TITLE OF INVENTION: MOLECULES, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Denver
STATE: Colorado
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                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 80203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         : 48 amino acids
amino acid
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1700 Lincoln Street, Suite 3500
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                                                                                                   2618-25-C1-1
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TYPE: amino acid;
TOPOLOGY: linear;
MOLECULE TYPE: protein
US-08-482-130C-65
US-08-906-769-65
; Sequence 65, Application US/08906769
; Patent No. 6077687
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Best Local Similarity
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                                                                                                                                                                 Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                       TELEFAX: (303) 863-02 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2618-25-C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 80203
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 48 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Frank, Glenn R.

APPLICANT: Stiegler, Gary L.

TITLE OF INVENTION: FLEA PROTEASE PROTEINS, NUC.

TITLE OF INVENTION: MOLECULES, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                252 VLIVLAVI 259
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Denver
STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/0:
FILING DATE: 07-JUN-1995
                                                                                                   1 VLIVLAVI 8
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STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                     (303) 863-0223
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100.0%; Pred. No.
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                      A;Cross-references: GB:AE000675; NID:g2982863; PIDN:AAC06501.1; A;Experimental source: strain VF5 C;Genetics: A;Gene: aq_142 C;Superfamily: Aquifex aeolicus hypothetical protein aq_142
                                                                                                                                                                                         Nature 392, 353-358, 1998
A;Title: The complete genome of the hyperthermophilic bacterium Aquifex aeolicus.
A;Reference number: A70300; MUID:98196666
A;Accession: F70313
                                                                                       A; Molecule type: DNA
A; Residues: 1-146 <AQF>
                                                                                                                A;Status:
                                                                                                                                                                                                                                                                                                                                                   preliminary; nucleic acid sequence
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PID:g2982881; GB:AE00

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conserved hypothetcial protein SA1250 [imported] - Staphylococcus aureus (strain C;Species: Staphylococcus aureus C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 22-Oct-2001 C;Accession: A99919
R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L. ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kalto, C.; Sekimiz C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001
Lancet 357, 1225-1240, 2001
Lincet 357, 1225-1240, 2001
Lincet 357, 1225-1240, 2001
A. Accession: A89758; MUID:21311952; PMID:11418146
 RESULT
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R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, Pham, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, ory, S.; Olson, M.V.
Tere 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                         type III export protein PscK PA1724 [imported] - Pseudomonas aeruginosa (strain C;Species: Pseudomonas aeruginosa C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000
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A; Residues: 1-208 <STO>
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A; Reference number: A82950;
A; Accession: D83431
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A;Experimental source: strain N315
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A; Residues: 1-204 <KUR>
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tive 0; Mismatches
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A; Description: Arabidopsis t
A; Reference number: Z14667
A; Accession: T02400
A.Grafin.
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A;Introns: 50/3; 74/1; 93/3; 119/1; 145/1; 174/2;
C;Superfamily: Agrobacterium beta-glucosidase
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A;Experimental source: Cultivar Columbia
A;Experimental source: Cultivar Columbia
R;Lin, X; Kaul, S; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon,
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon,
M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Fraser, C.M.; Venter
M.; Koo, H.; Moffat, K.S.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter
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C;Superfamily: Arabidopsis thaliana hypothetical
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C;Accession: E85276
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A;Molecule type: DNA
A;Residues: 1-506 <ROUD
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A; Residues: 1-959 <STO>
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Nature 402, 761-768, 1999
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Library, May 1998
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G3P3_YEAST
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G3P1_TRIKO
PTWC_ECOLI
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CSD_BORBU
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YS81_CAEEL
CP84_RABIT
CP85_RABIT
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2 escherichia
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ALIGNMENTS

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EXSB_PSEAE P26994;
                                                                                                                                                                               TRANSMEM
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                  "The organization and sequence of the genes for ATP synthase subunits in the cyanobacterium Synechococus 6301. Support for an endosymbiotic origin of chloroplasts.";

J. Mol. Biol. 194:359-383(1987).

1- FUNCTION: A POSSIBLE FUNCTION FOR THIS PROTEIN IS TO GUIDE THE ASSEMBLY OF THE MEMBRANE SECTOR OF THE ATPASE ENZYME COMPLEX.

-i- SIMILARITY: TO THE CORRESPONDING SUBUNIT IN OTHER BACTERIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE-87311713; PubMed-
Cozens A.L., Walker J.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AE000675; AAC06501.1; -. Hypothetical protein; Complete SEQUENCE 146 AA; 16321 MW;
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PIR; S10825; (
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SEQUENCE FROM N.A.

(STRAIN-ATCC 15692 / PAO1;

(MEDLINE-20437337; PubMed-10984043;

X MEDLINE-20437337; PubMed-10984043;

X Hickey M.J., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagrova Garber R.L., Goltry L., Tolentino E., Westbrock-Wadman S., Yuan Y., Brody L.L., Coulter S.N., Folger K.R., Kas A., Larbig K., Lim R. Smith K.A., Spencer D.H., Wong G.K.-S., Wu Z., Paulsen I.T., A Smith K.A., Saier M.H., Hancock R.E.W., Lory S., Olson M.V.;

"Complete genome sequence of Pseudomonas aeruginosa PAO1, an opportunistic pathogen.";

"I "ature 406:959-964(2000).

"I "SIMILARITY: TO Y.ENTEROCCLITICA VIRB.
                                                                                                                                                                                                                                                                         Query
Best L
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STRAIN-ATCC 15692 / PAO1;
STRAIN-ATCC 15692 / PAO1;
MEDIJINE-92011420; PubMed-1655713;
Frank D.W., Iglewski B.H.;
"Cloning and sequence analysis of a trans-regulatory locus required for excenzyme S synthesis in Pseudomonas aeruginosa.";
""""reriol. 173:6460-6468(1991).
                                                                                                                                                                                                                                                                                                                    EMBL; AE004597; AAG05
PIR; B41047; B41047.
Complete proteome.
CONFLICT 62
SEQUENCE 137 AA; 1
                                                                               30-MAY-2000 (Rel. 39, 1
16-OCT-2001 (Rel. 40, 1
X-linked juvenile retiners) OR RS1 OR RS1H OR XLRS1.
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16-OCT-2001
Exoenzyme S
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                                            Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Bacteria; Proteobacteria;
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SEQUENCE FROM N.A
STRAIN=129/SVJ;
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length: 2000000000
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O9ddBf7 mus musculu
O9df11 homo sapien
O9bx67 homo sapien
O9bx67 homo sapien
O99u70 staphylococ
P95439 pseudomonas
O9i313 pseudomonas
O9i313 pseudomonas
O9i313 pseudomonas
O9i446 ctenocephal
O9i144 arabidopsis
O64879 arabidopsis
O62987 arabidopsis
O99gx8 heliocoverp
O9fe08 perilla fru
O93hq4 streptococc
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ALIGNMENTS

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JCAM2 OR JAM	JUNCTIONAL A	01-DEC-2001	01-MAR-2001	01-MAR-2001	Q9EPK4;	Q9EPK4	K4	LT 1
JCAM2 OR JAM-2 OR 1110002N23RIK.	ADHESION MOLECUL	(TrEMBLrel. 19,	(TrEMBLrel. 16,	(TrEMBLrel. 16,		PRELIMINARY;		
23RIK.	JUNCTIONAL ADHESION MOLECULE-2, JAM-2 (1110002N23RIK PROTEIN	01-DEC-2001 (TrEMBLrel. 19, Last annotation update)	Last sequence update)	Created)		PRT; 310 AA.		
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Kondo S., Yamanaka I.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Schriml L.M., Staubii F., Szuki R., Tomita M., Wagner L., Washio T.,
RA Schriml L.M., Staubii F., Szuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Wordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mature 409:685-690(2001).

EMBL; AJ300304; CAC20704.1; -.

EMBL; AK013156; BAB28683.1; -.

MGD; MGI:1933820; Jcam2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=C57BL/6J; TISSUE=EMBRYO;
MEDLINE=21085660; PubMed=11217851;
298
                                                                                                                                                                                                                                                                                                                                                                                                                                   SMART;
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InterPro; IPR003600; Ig_like.
InterPro; IPR003006; Ig_MHC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Functional annotation
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XT; SM00409; IG; 2.
XT; SM00408; IGc2; 2.
XT; SM00410; IG_like; 1.
                                                                                                                                                                                       VALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRND
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EEGDFRHKSSFVI
                             EEGDFRHKSSFVI 310
                                                                                                                            VPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEV
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100.0%;
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1 JAM-3: an
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Pred. No. 1.5e-252;
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O9D1M9;
01-JUN-2001 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                         InterPro; IPR003006; Ig_MHC.
Pfam; PF00047; ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00408; IGc2; 2.
SMART; SM00410; IG_like; 1.
Immunoglobulin domain.
SEQUENCE 310 AA; 34819 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
STRAIN=C57BL/6J; TISSUE=EMBRYO;
MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Functional annotation of a full-length mouse Nature 409:685-690(2001).
EMBL; AK003326; BAB22715.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kawai J., Shinagawa A., Shibata
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InterPro; IPR003600; Ig_like
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VALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRND
                                                                        VPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCIASNDAGAARCEGQDMEV
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                                                                                                                 VALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRND
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                                                         VPLPTDSRANPRFQNSSFHVNSETGTLVFNAVHKDDSGQYYCTASNDAGAARCEGQDMEV
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Last sequence update)
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Pred. No. 2.8e-244;
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuchi P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubii F., Suzuki R., Tomita M., Wagner L., Washio T.,
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RA Sakai K., Okido T., Frunno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Buit C., Fletcher C., Fujita M., Gariboldi M.,
RA Basaki H., Sato K., Schoenbach C., Seya T., Kamiya M., Lee N.H.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPace Process of the Pfam; PF0004, Pfam; PF000409; IG, SMART; SM00408; IGC2; 2.
SMART; SM00410; IG_like; 1.
SMART; SM00410 id_like; 1.
310 AA; 34855 MW; C
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01-JUN-2001 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLRel. 1
1110002N23RIK PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Functional annotation of a full-length mouse cDNA collection."; Nature 409:685-690(2001).
Nature 409:685-690(2001).
EMBL; AK008187; BAB25519.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        298 EEGDF
                                                                                                                                                                                                                                                                                               58
                                                                                                                                                                                                                                                                                                                                                  58
                                                      VPLPTDSRANPREQNSSEHVNSETGTLVENAVHKDDSGQYYCIASNDAGAARCEGQDMEV
                                                                                                                                                                                                                                                                                                                       SQTSDPRIEWKKIQDGQTTYYYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEV 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MGI:1933820; Jcam2.
YDLNIAGIIGGVLVVLIVLAVITMGICCAYRRGCFISSKQDGESYKSPGKHDGVNYIRTS
                                                                                                                                                                                                    VALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYSWYRND
                                                                                                                                                                      VALNDRKEVDEITIELIVQVKPVTPVCRIPAAVPVGKTATLQCQESEGYPRPHYNWYRND
                                                                                                                                                                                                                                                                                            SQTSDPRIEWKKIQDGQTTYVYFDNKIQGDLAGRTDVFGKTSLRIWNVTRSDSAIYRCEV
                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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IPR003600;
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                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                          49.0%;
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Ig_like.
Ig_MHC.
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17,
19,
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Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                           Score 152; DB Pred. No. 2.6e 0; Mismatches
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C74884EABE234680 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 11;
.6e-148;
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; Murinae; Mus
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297
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RESULT
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Best Local S
Matches 24
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Q96FL1;
01-DEC-2001
01-DEC-2001
01-DEC-2001
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junctions.";
Submitted (AUG-2001) to the
EMBL; AF356518; AAK27221.1;
                                                                                        expression and
                                                                                                                                     TISSUE=BRAIN;
Cunningham S.A., Arrate M.P., Tran T.M.;
"Cloning of Human Junctional Adhesion Molecule
"Cloning of Human Junctional Adhesion Molecule
Submitted (MAR-2001) to the EMBL/GenBank/DDBJ d
                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE-EYE, AND Strausberg R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-DEC-2001 (TYEMBLrel. 19, Created)
01-DEC-2001 (TYEMBLrel. 19, Last sequence update)
01-DEC-2001 (TYEMBLrel. 19, Last annotation update)
UNKNOWN (PROTEIN FOR IMAGE:3875338) (FRAGMENT).
                                                                   family members.";
Submitted (AUG-2001)
                                                                                                 Aurrand-Lions M.A., Johnson-leger C., Wong C., DuPasquier 
"Heterogeneity of endothelial junctions is reflected by d
                                                                                                          SEQUENCE FROM N.A.,
Aurrand-Lions M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                           Aurrand-Lions
                                                   SEQUENCE FROM
                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9606;
                                Junctional adhesion
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                                                                                                                                                                                                                                                                                                                                                      PRPHYSWYRNDVPLPTDSRANPRF
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EEGDFRHKSSFVI 310
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                                                                                                                                                                                                                                                                                                                                                                                                                                        309 AA;
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Human).
                                         M.A.
                                                                                 specific
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                                                                                                                                                                                                         Chordata;
Primates;
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Primates;
                                         Johnson-leger
                                                                                                                                                                                                                                                                                                                                                                                                                                        34917 MW;
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              EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                  Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                      189
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                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 3. 
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Craniata; Ve
Catarrhini;
                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                               ger C.,
(JAMS)
                                                                                                                                                                                                                                                                                                                                                                                                                                        50C5B1B7872E8DF3 CRC64
                                                                                       localization
                                Lamagna
and inte
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i; Hominidae; Homo.
                                                                                                                                                                                                                                                       ion update)
(JUNCTIONAL
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                                 interendothelial
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                                           Ozaki
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differential
                                                                                        three
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ID P95439;
AC P95439;
                                                                RESULT \7
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C Q99U70;

O1-JUN-2001 (TrEMBLrel. 17, Created)

O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)

E CONSERVED HYPOTHETCIAL PROTEIN (HYPOTHETICAL PROTEIN SAV1417).

S Staphylococcus aureus (strain N315), and

S Staphylococcus aureus (strain Mu50).

C Bacillus/Staphylococcus group;

C Bacillus/Staphylococcus group;

C Bacillus/Staphylococcus group;

C Bacillus/Staphylococcus group;

C NCBI_TaxID-158879, 158878;

N [1]
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Best Local S
Matches 8
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Best Local Similarity
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Pfam; PF00047; 1g; 2.
SMART; SM00409; IG; 2.
SMART; SM00408; IGc2; 2.
SMART; SM00410; IG_like; 1.
Immunoglobulin domain; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lancet 357:1225-1240(2001).
EMBL; AP003133; BAB42510.1; -.
EMBL; AP003362; BAB57579.1; -.
Interpro; IPR00326; PA_PTPase.
Pfam; PF01569; PAP2; 1.
SMART; SM00014; acidPPc; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.

SPECIES-S.aureus (strain N315), and S.aureus (strain Mu50);

MEDLINE-21311952; PubMed-11418146;

Kuroda M., Ohta T., Uchiyama I., Baba T., Yuzawa H., Kobayashi I
Cui L., Oguchi A., Aoki K.-I., Nagai Y., Lian J.-Q., Ito T.,
Kanamori M., Matsumaru H., Maruyama A., Murakami H., Hosoyama A.
Mizutani-Ui Y., Takahashi N.K., Sawano T., Inoue R.-I., Kalto C.
Sekimizu K., Hirakawa H., Kuhara S., Goto S., Yabuzaki J.,
Kanehisa M., Yamashita A., Oshima K., Furuya K., Yoshino C., Shi
Hattori M., Ogasawara N., Hayashi H., Hiramatsu K.;
"Whole genome sequencing of meticillin-resistant Staphylococcus
aureus.";
                                                                                                                                                                                                                                                                                                                                                                                                        Complete SEQUENCE
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InterPro;
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                                                                                                                                                                                                                       242 IAGIIGGV 249
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rPro; IPR003599;
rPro; IPR003598;
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                                                                                                                                                                                                                                                                          Similarity
8; Conser
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204 AA;
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llarity 100.0%;
Conservative
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llarity 100.0%;
Conservative
                                           PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein.
22685 MW; 009CF65D8B1FBDF2 CRC64;
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; Ig_like.
; Ig_MHC.
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                                                                                                                                                                                                                                                                                                     Score 8;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    POTENTIAL.
; CE39ADF33EA1DAB9 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                            Mismatches
                                           206 AA
                                                                                                                                                                                                                                                                                                     DB 16;
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hes 0;
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Kaito C.,
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STRAIN-ARCC 15692 / PAO1;

MEDLINE-20437337; PubMed-10984043;

MEDLINE-204437337; PubMed-10984043;

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Hickey M.J., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagrou M.

Garber R.L., Goltry L., Tolentino E., Westbrock-Wadman S., Yuan Y.,

Brody L.L., Coulter S.N., Folger K.R., Kas A., Larbig K., Lim R.M.,

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                                                                                                                                                                                                     EMBL; AE004598; AAG05113.1;
Complete proteome.
SEQUENCE 208 AA; 23898 MV
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Yahr T.L., Goranson J., Frank D.W.;
"Exoenzyme S of Pseudomonas aeruginosa
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Bacteria; Proteobacteria;
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EMBL; U56077; AAC44782.1; -.
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